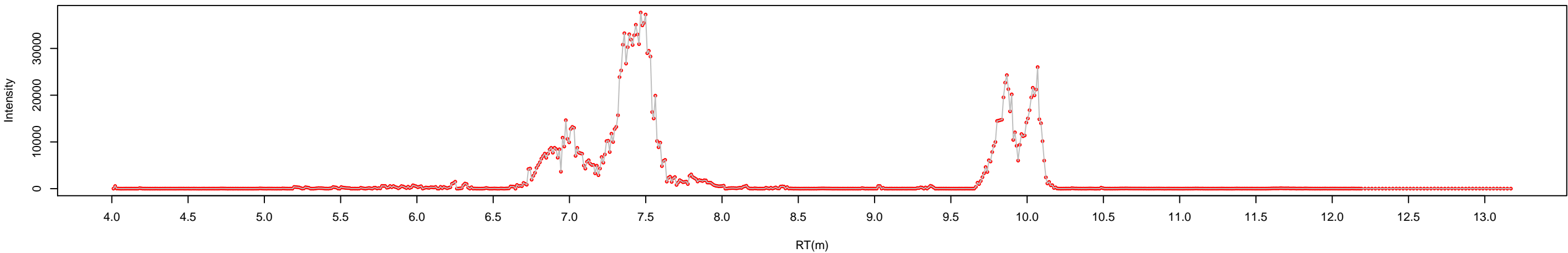
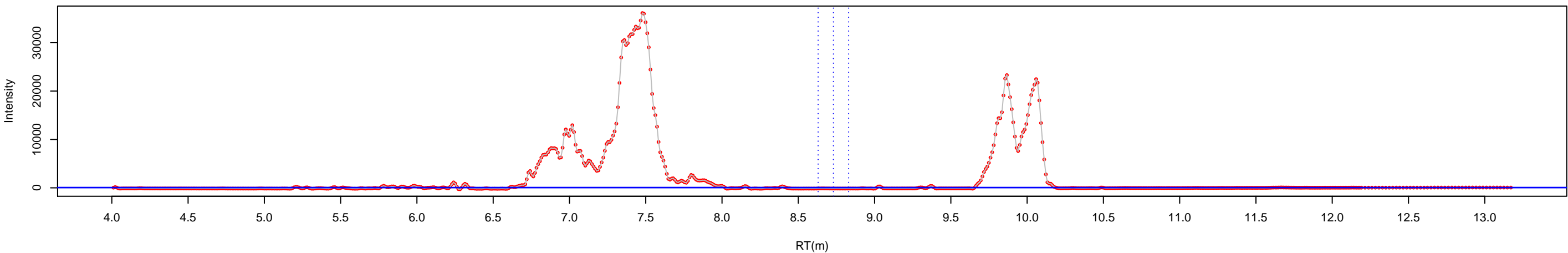


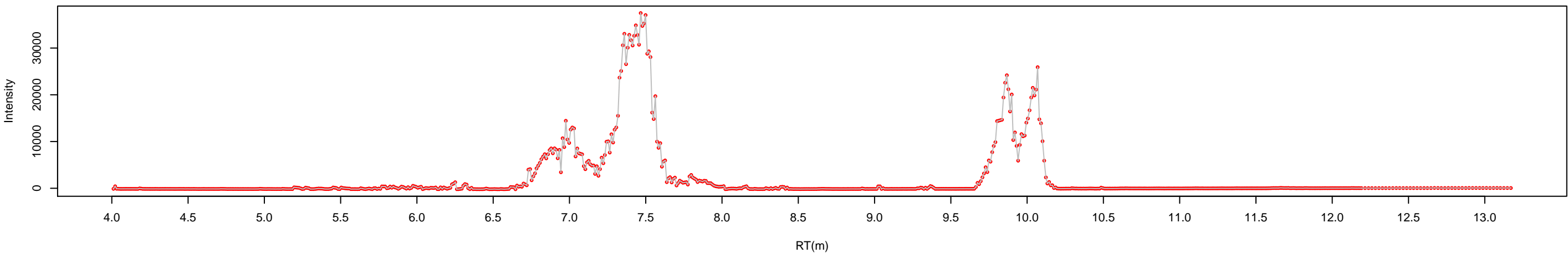
raw chromatogram | batch: 1 sample: coth10 conc: NA function: 2 mass: 514.35>79.93



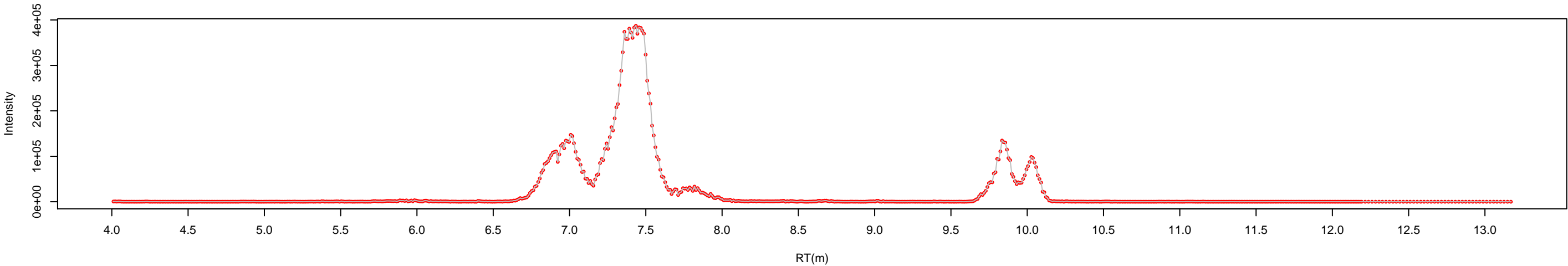
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 9.73 BLine: yes



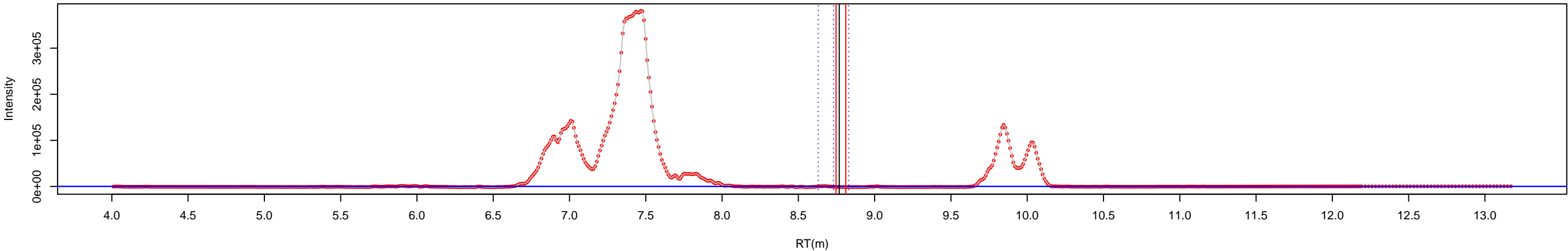
peak area | window size:3 BLine: yes



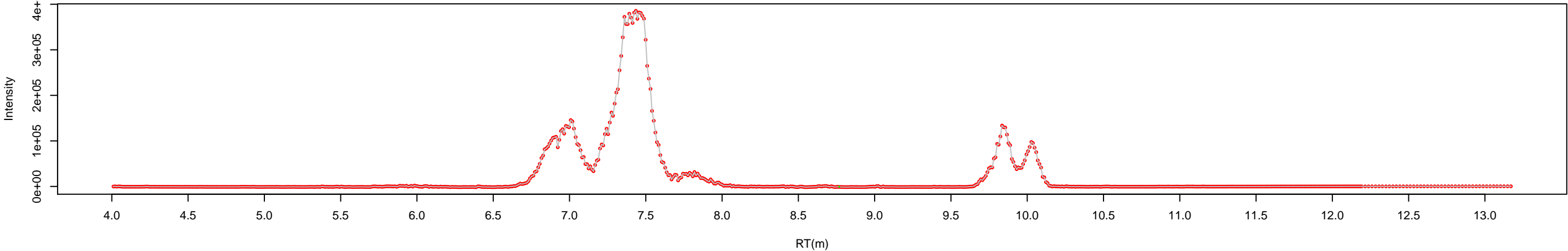
raw chromatogram | batch: 1 sample: cotn19 conc: NA function: 2 mass: 514.35>79.93



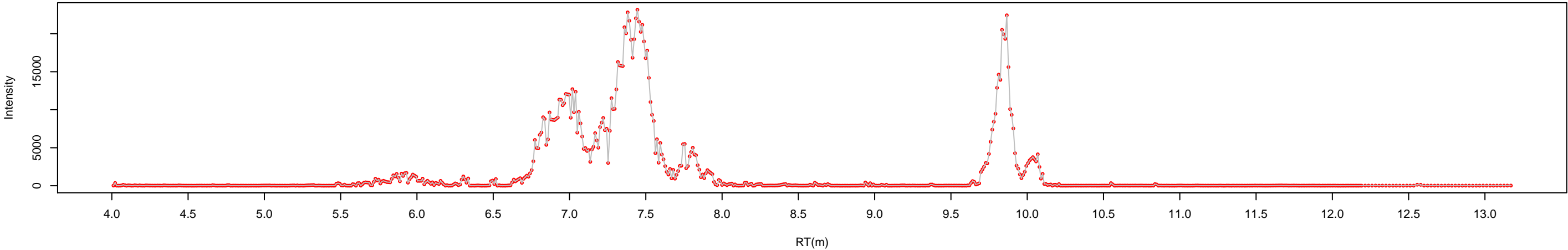
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 17.6 BLine: yes



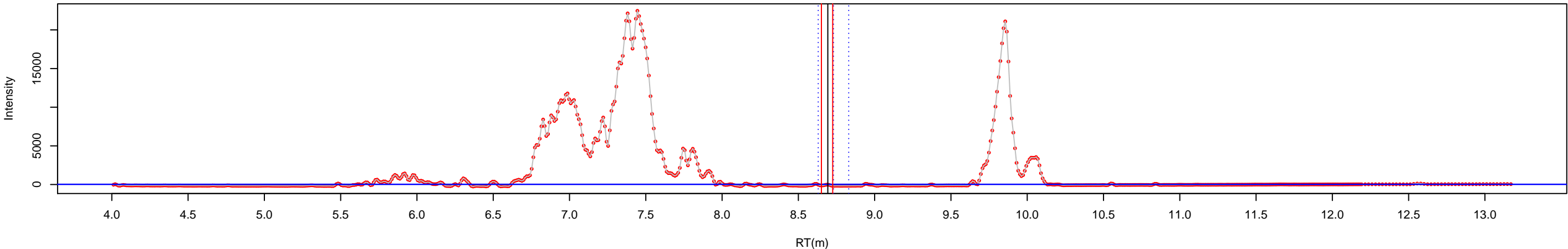
peak area | window size:3 BLine: yes



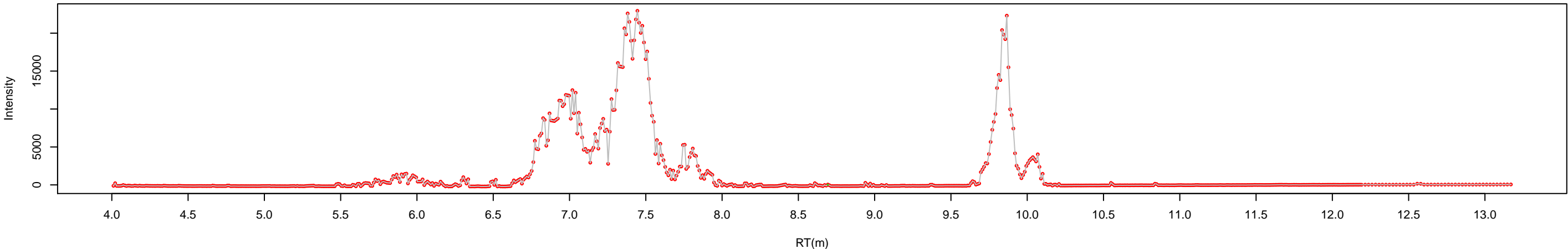
raw chromatogram | batch: 1 sample: coth11 conc: NA function: 2 mass: 514.35>79.93



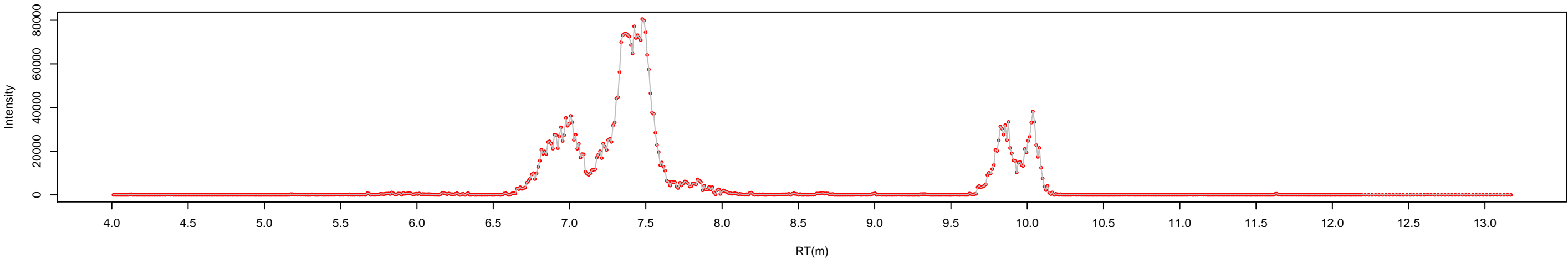
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 4.18 BLine: yes



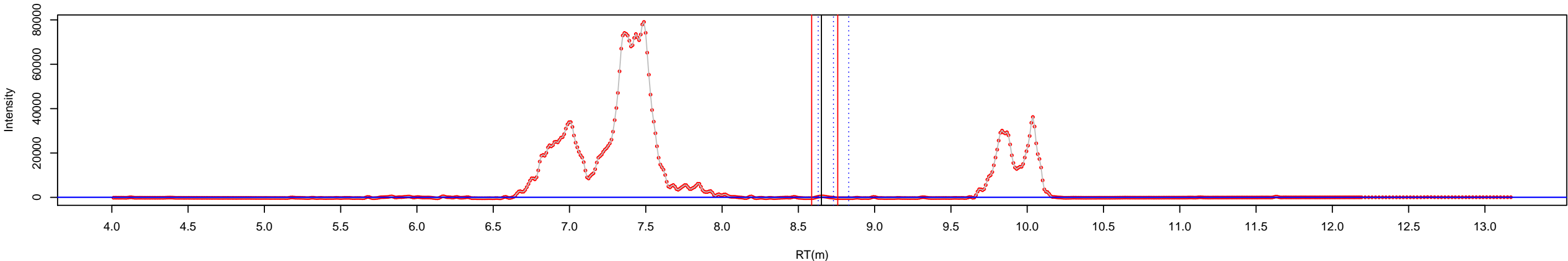
peak area | window size:3 BLine: yes



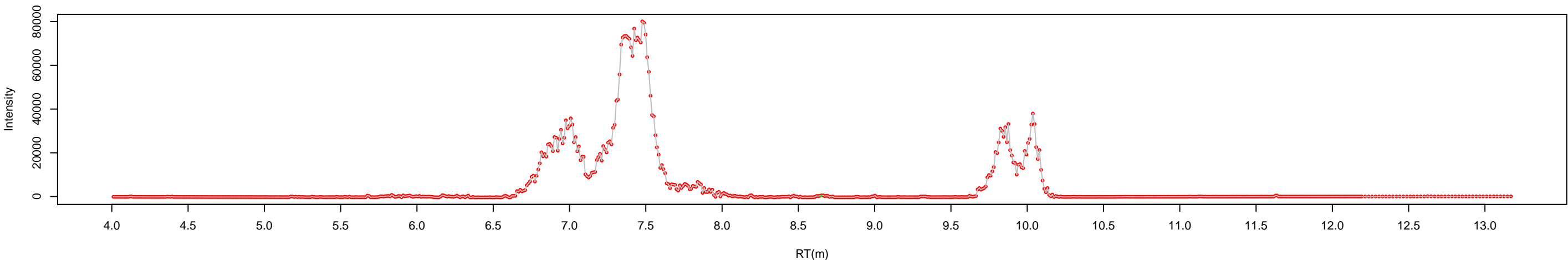
raw chromatogram | batch: 1 sample: cotn20 conc: NA function: 2 mass: 514.35>79.93



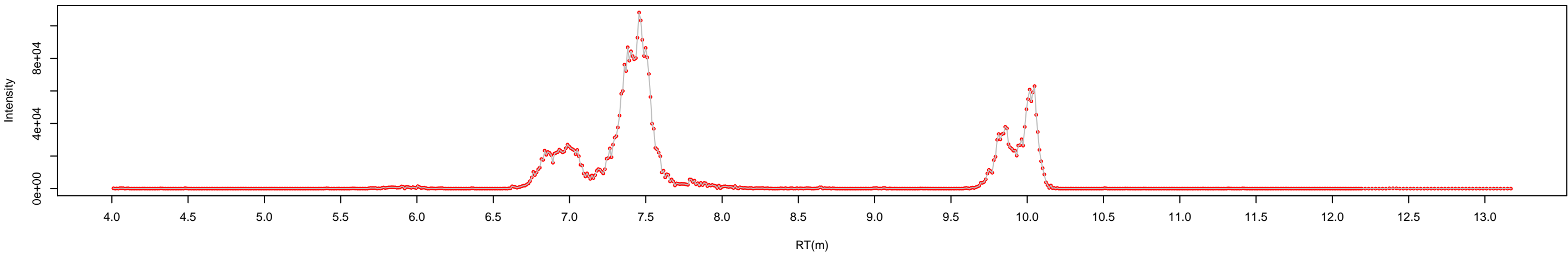
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 7.56 BLine: yes



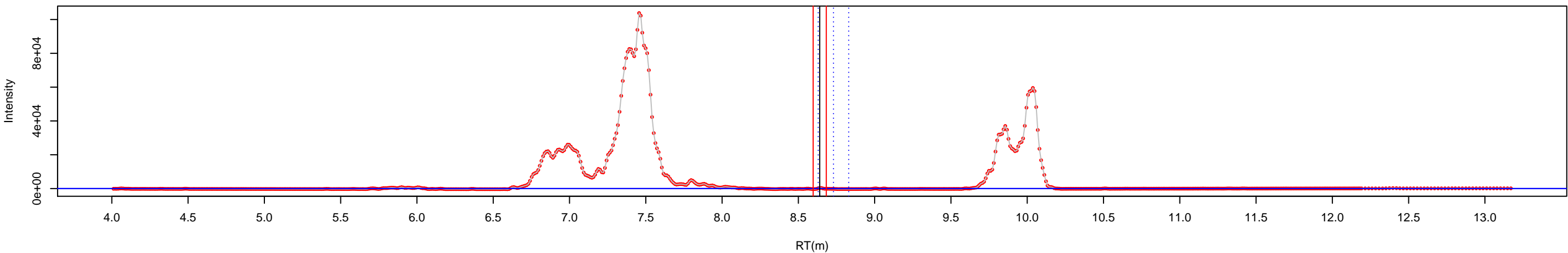
peak area | window size:3 BLine: yes



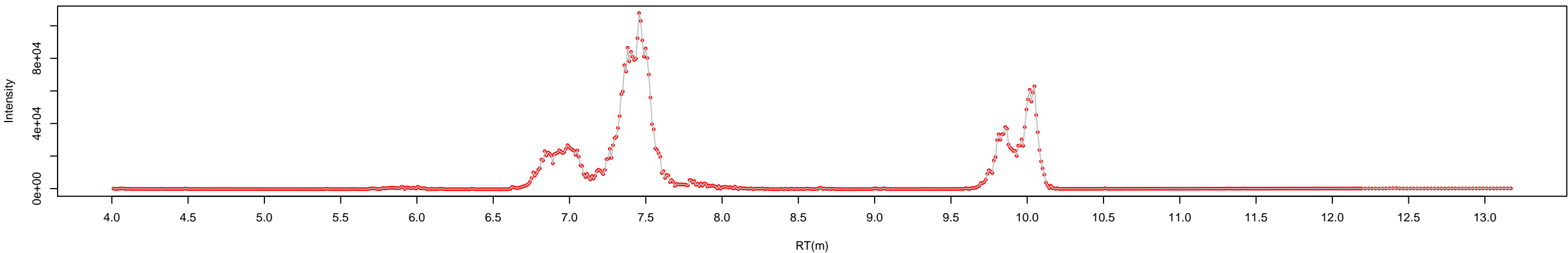
raw chromatogram | batch: 1 sample: coth12 conc: NA function: 2 mass: 514.35>79.93



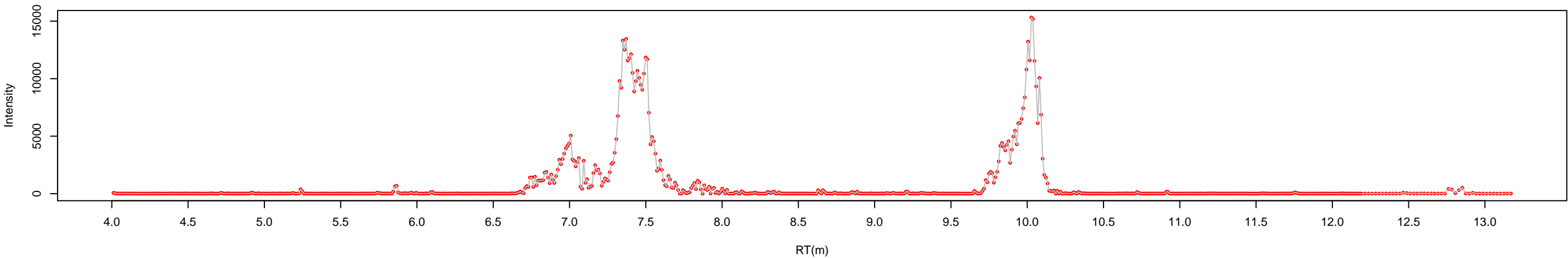
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 3.46 BLine: yes



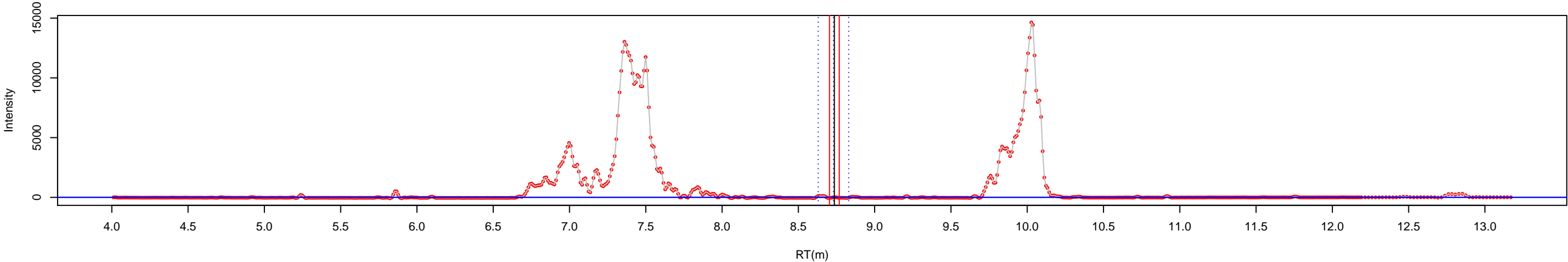
peak area | window size:3 BLine: yes



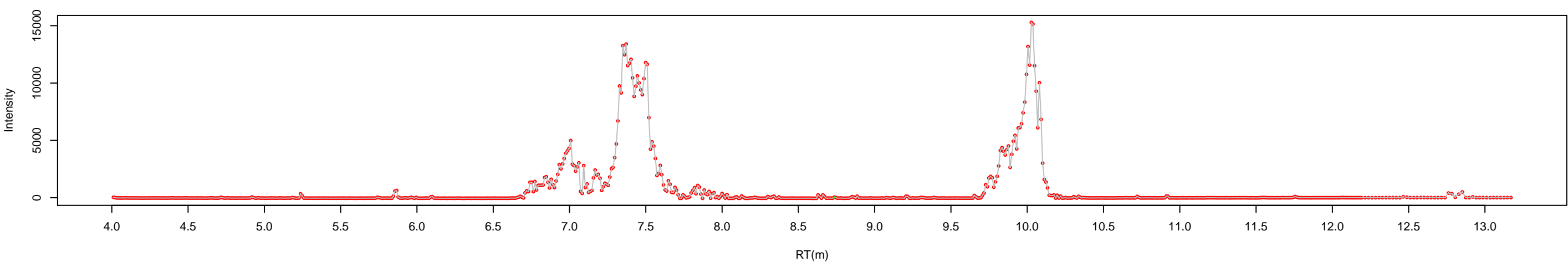
raw chromatogram | batch: 1 sample: cotn21 conc: NA function: 2 mass: 514.35>79.93



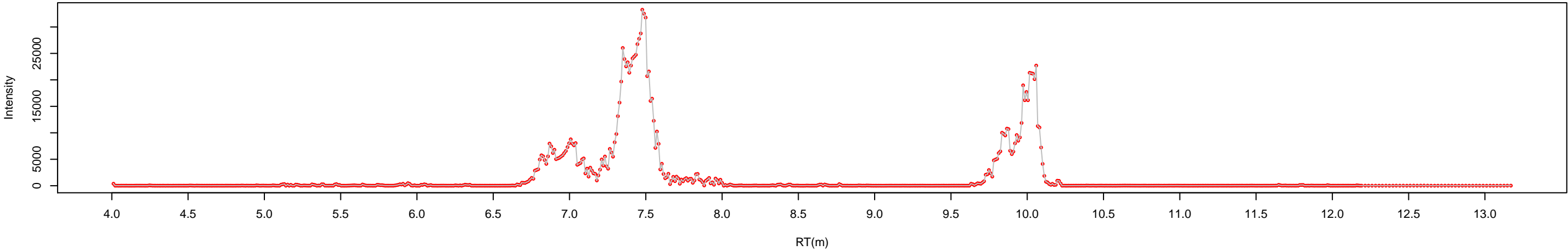
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.674 BLine: yes



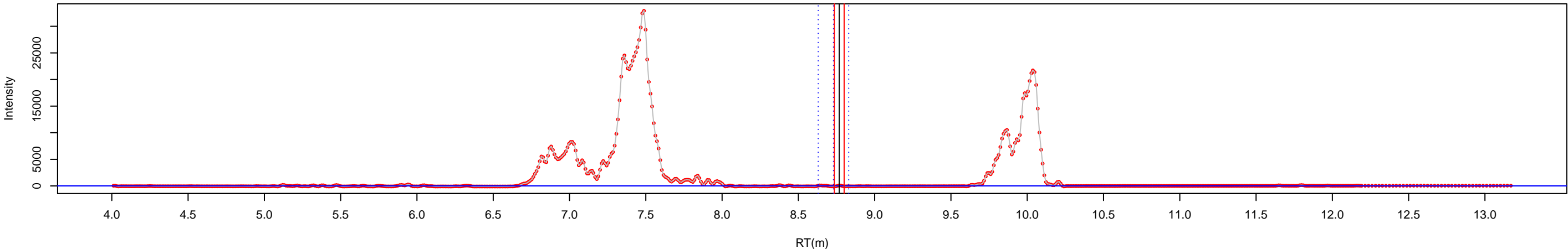
peak area | window size:3 BLine: yes



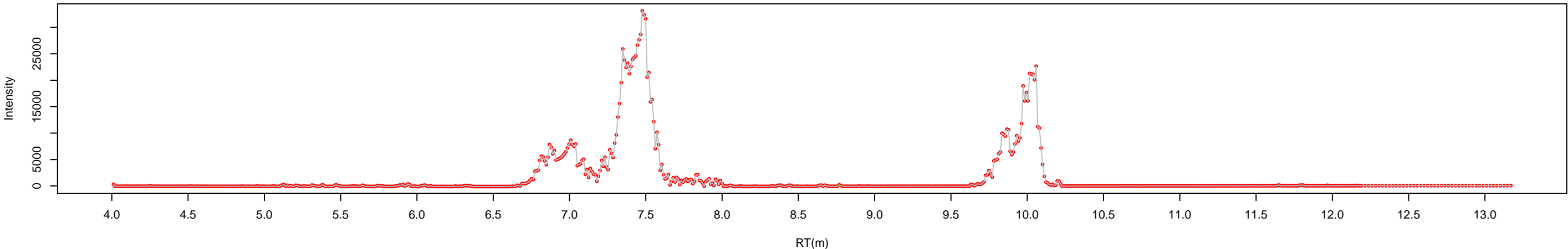
raw chromatogram | batch: 1 sample: coth13 conc: NA function: 2 mass: 514.35>79.93



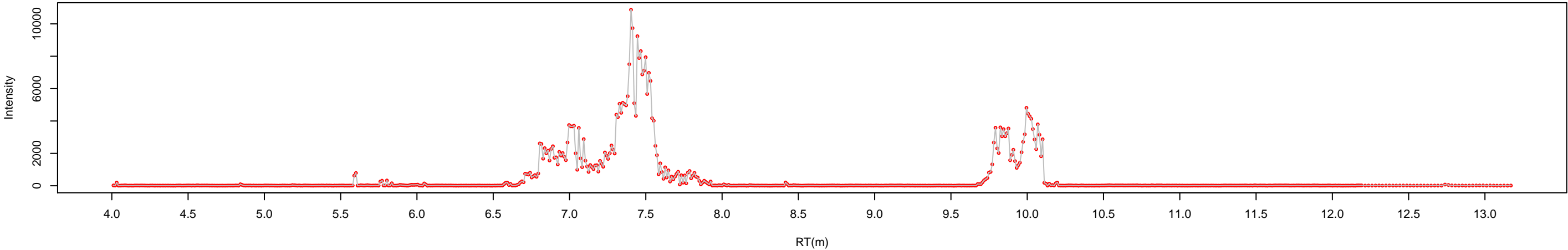
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 2.56 BLine: yes



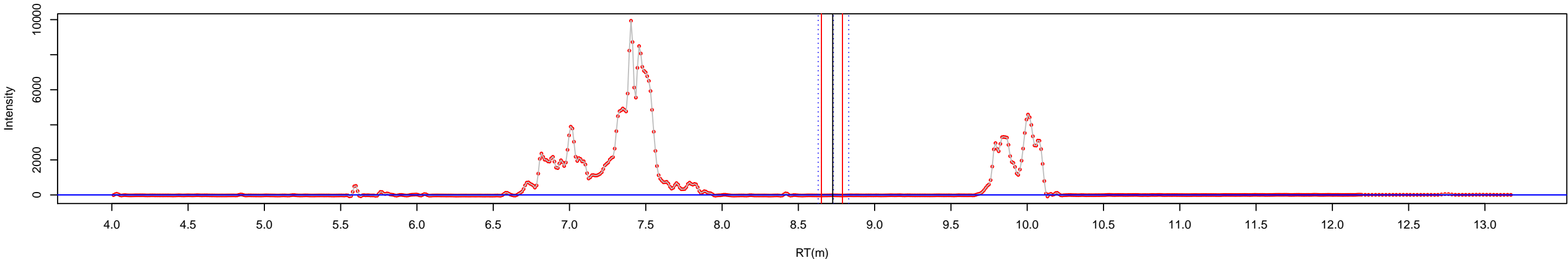
peak area | window size:3 BLine: yes



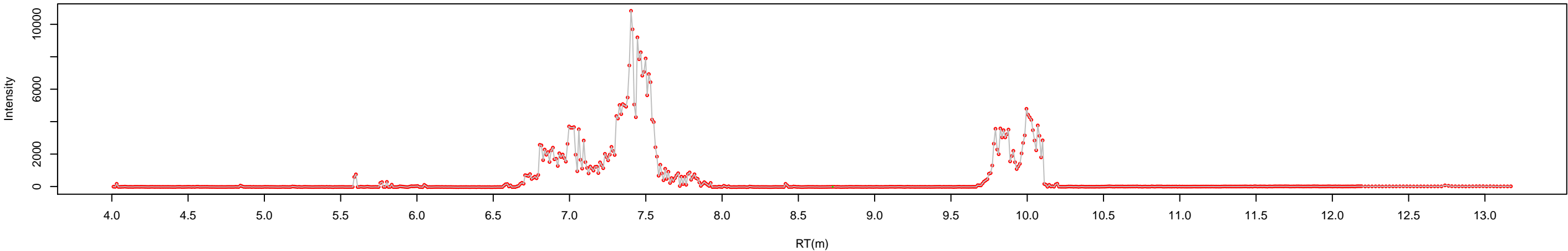
raw chromatogram | batch: 1 sample: cotn22 conc: NA function: 2 mass: 514.35>79.93



peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.436 BLine: yes

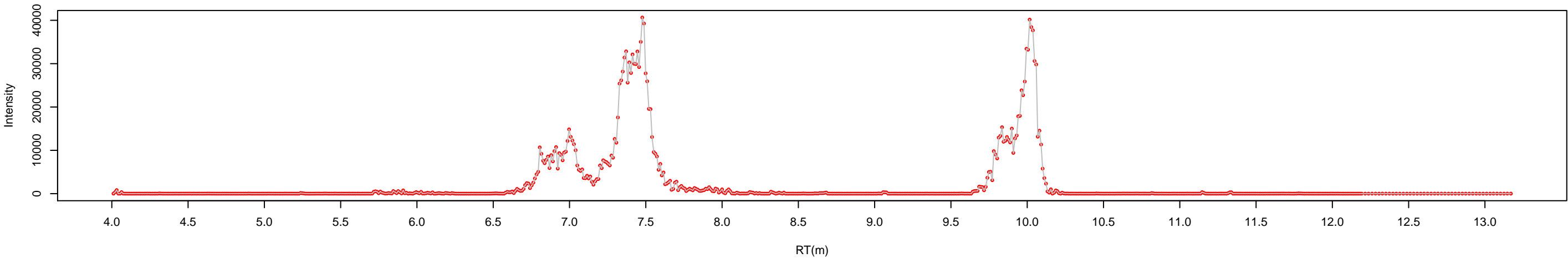


peak area | window size:3 BLine: yes

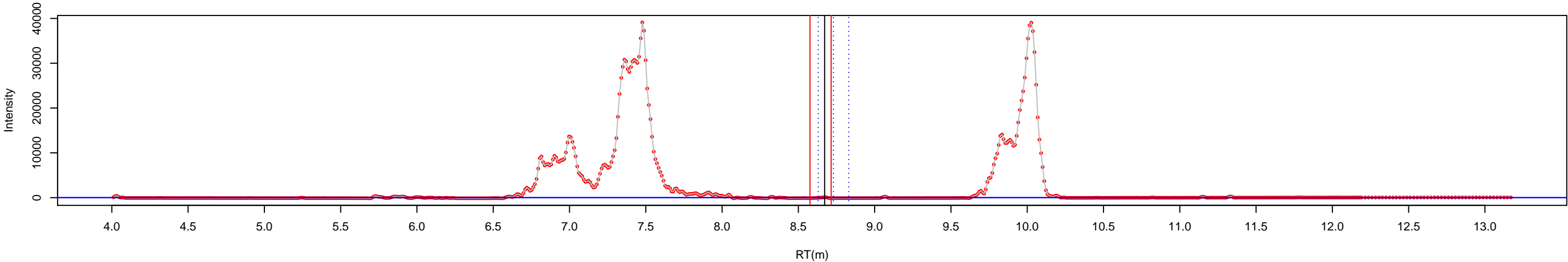




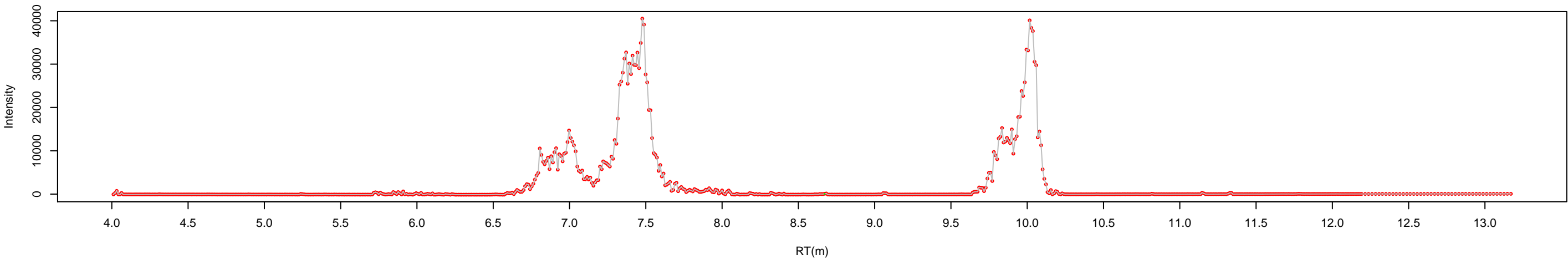
raw chromatogram | batch: 1 sample: coth14 conc: NA function: 2 mass: 514.35>79.93



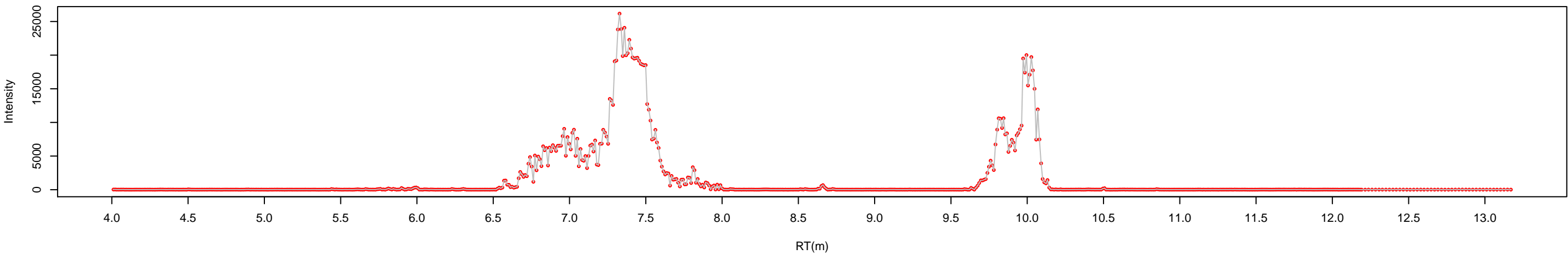
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 1.97 BLine: yes



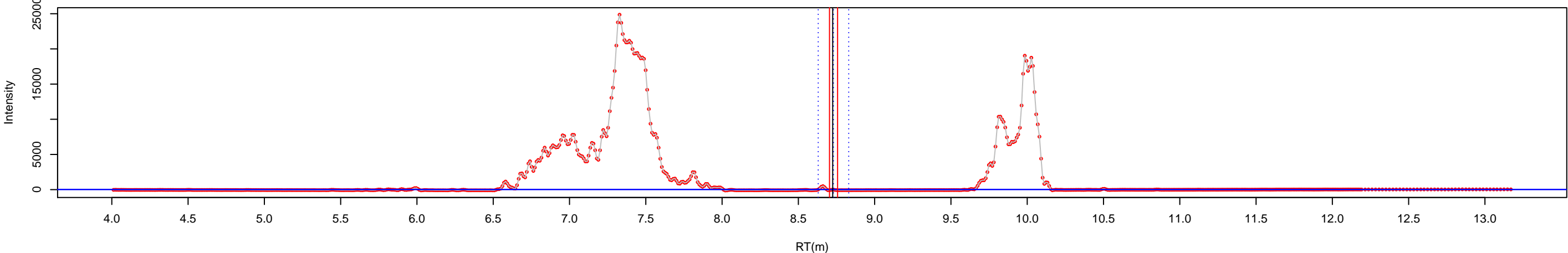
peak area | window size:3 BLine: yes



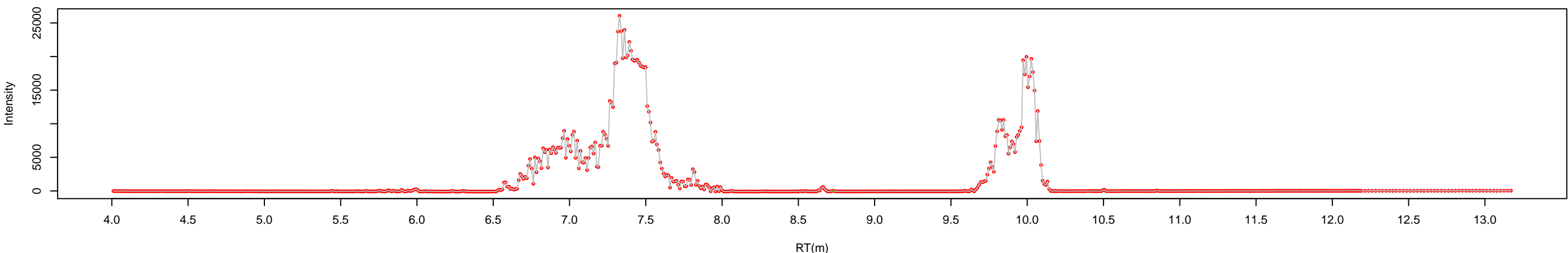
raw chromatogram | batch: 1 sample: cotn23 conc: NA function: 2 mass: 514.35>79.93



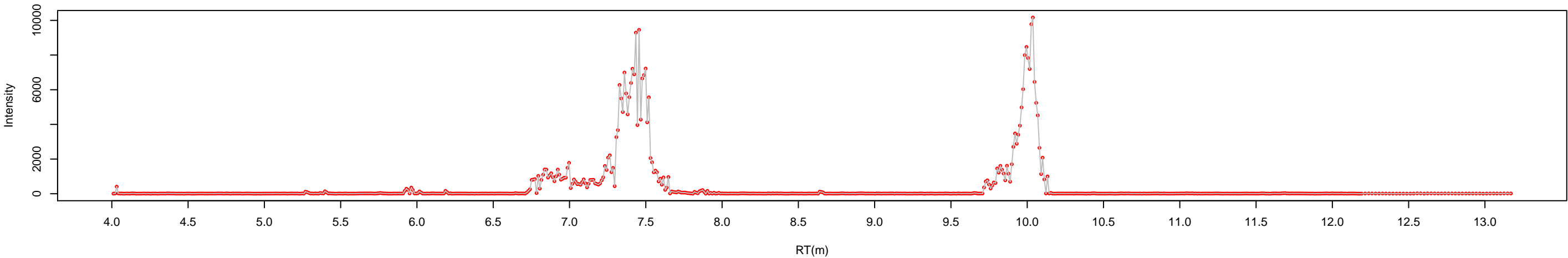
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 3.53 BLine: yes



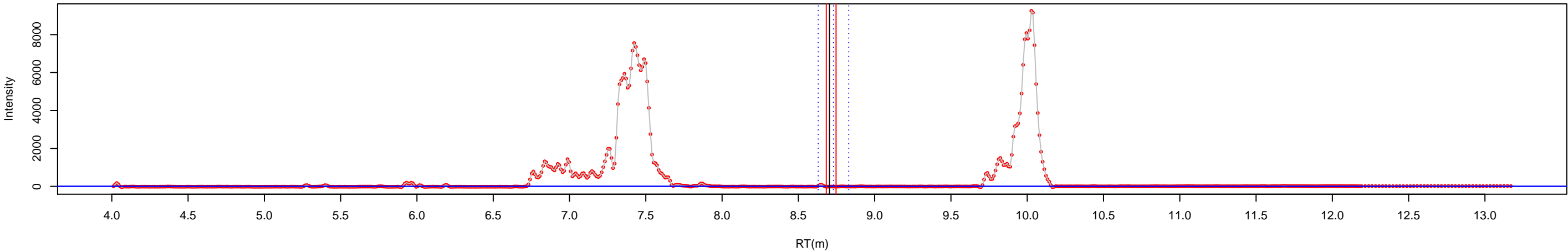
peak area | window size:3 BLine: yes



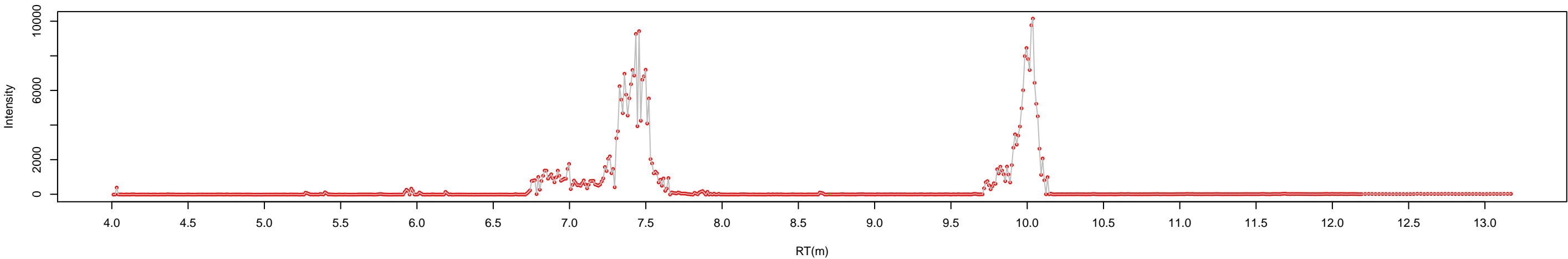
raw chromatogram | batch: 1 sample: coth15 conc: NA function: 2 mass: 514.35>79.93



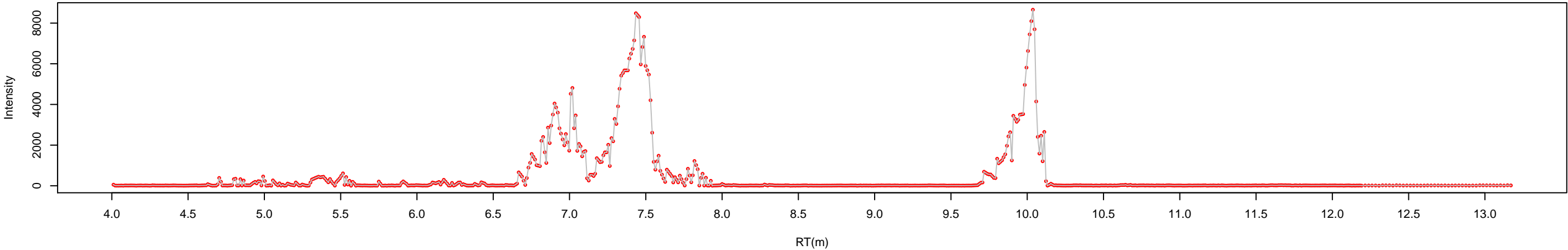
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.433 BLine: yes



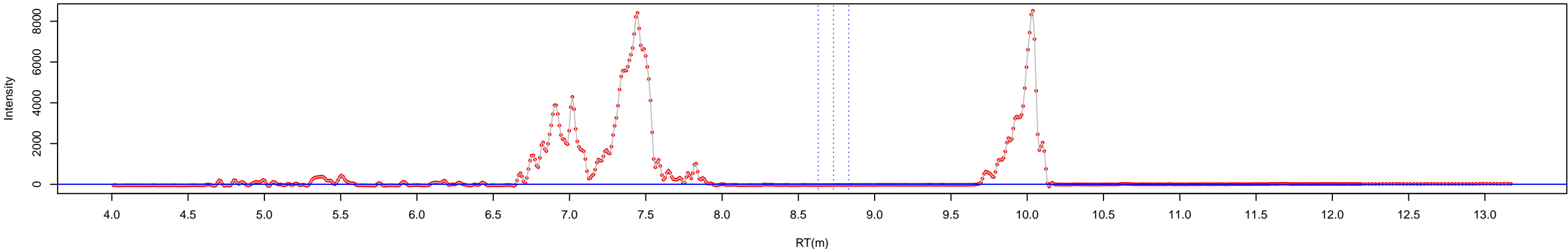
peak area | window size:3 BLine: yes



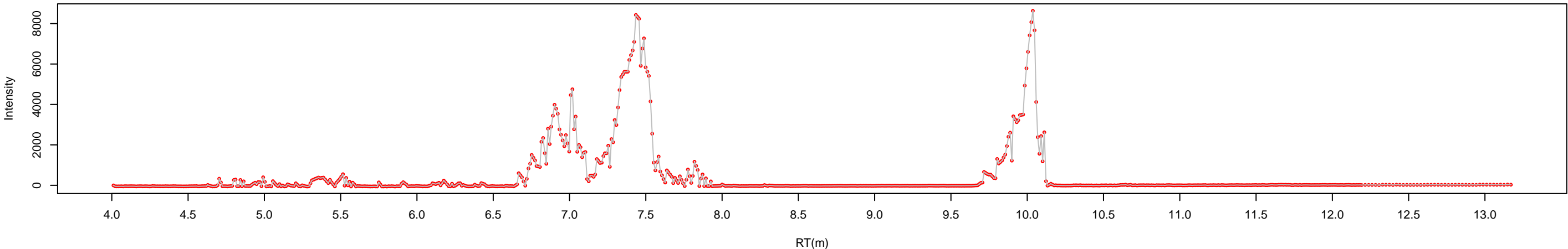
raw chromatogram | batch: 1 sample: cotn24 conc: NA function: 2 mass: 514.35>79.93



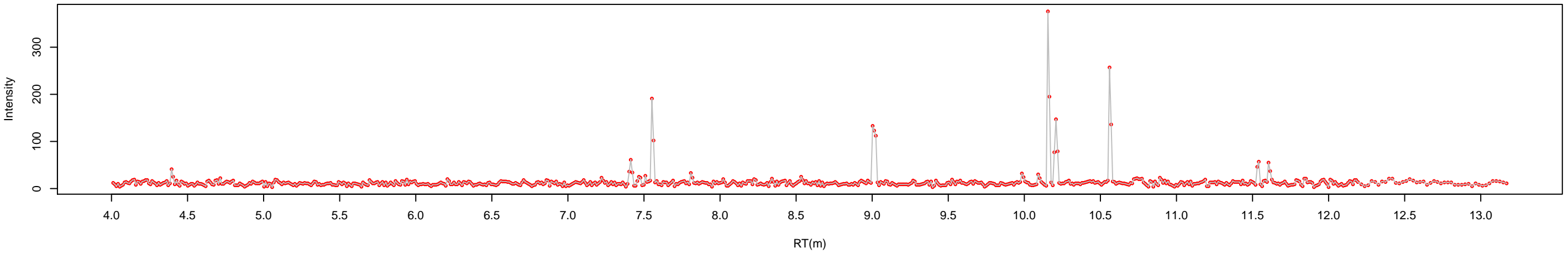
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.716 BLine: yes



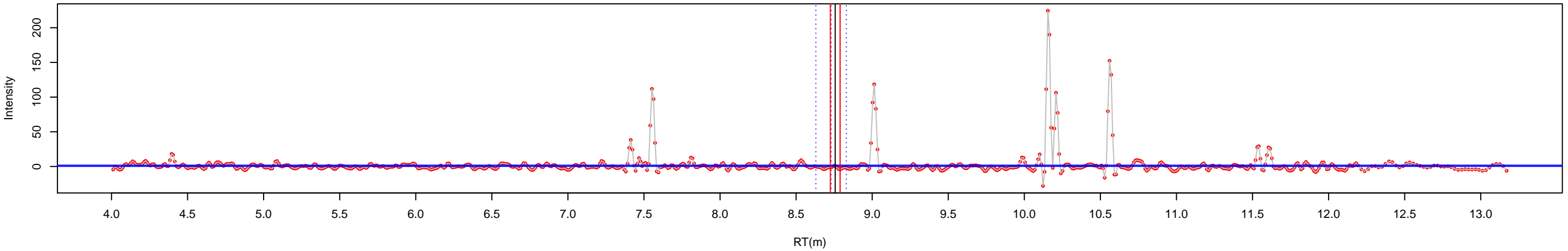
peak area | window size:3 BLine: yes



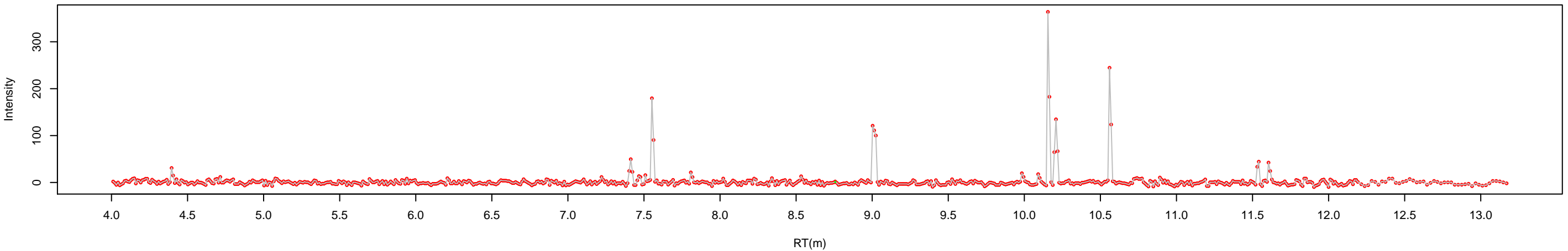
raw chromatogram | batch: 1 sample: std0-10ul-1 conc: 0.001 function: 2 mass: 514.35>79.93



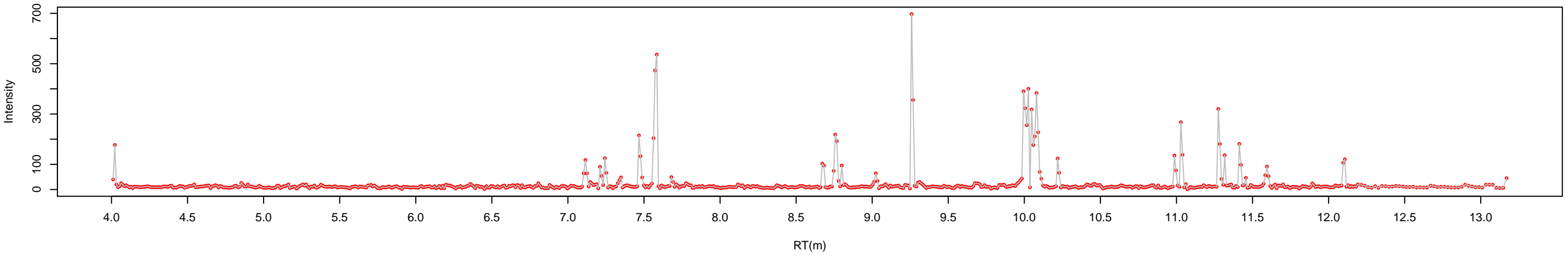
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.177 BLine: yes



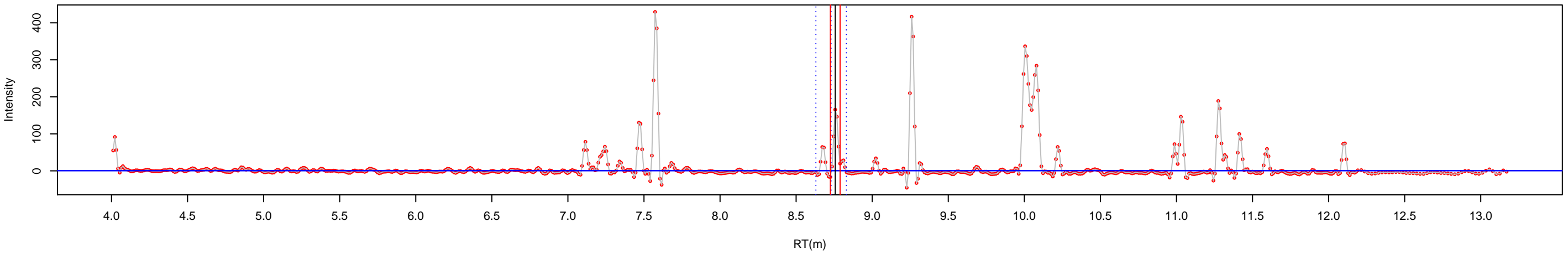
peak area | window size:3 BLine: yes



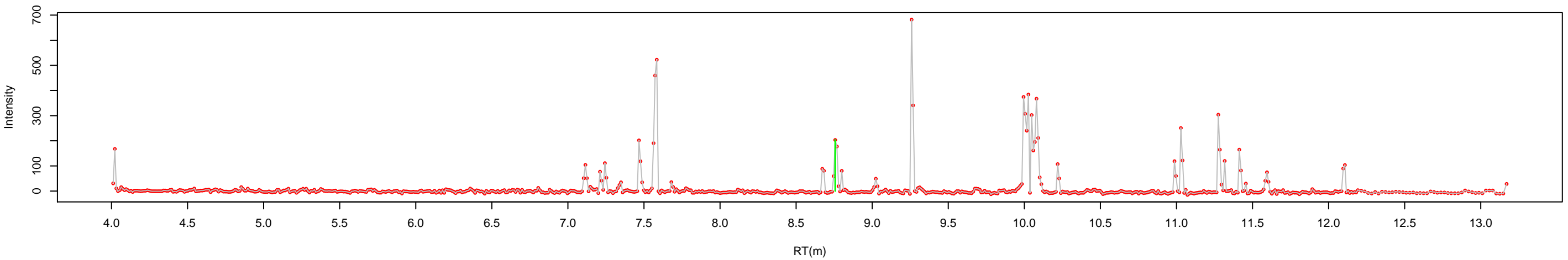
raw chromatogram | batch: 1 sample: std1-10ul-1 conc: 0.002 function: 2 mass: 514.35&gt;79.93



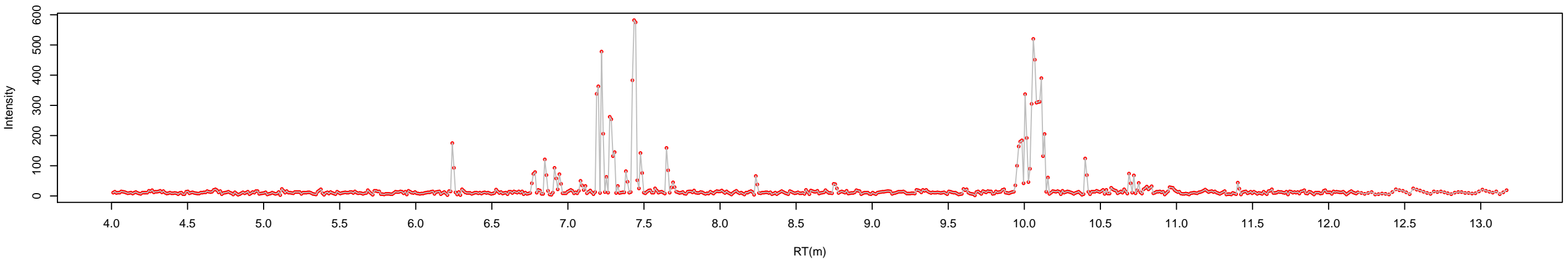
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.106 BLine: yes



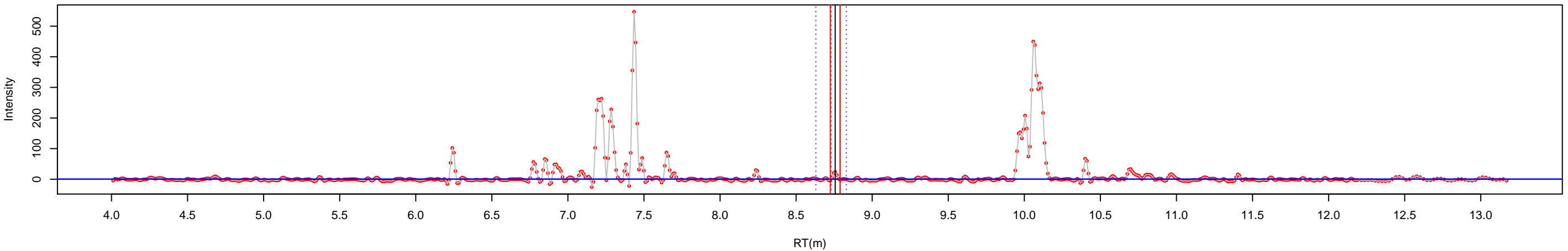
peak area | window size:3 BLine: yes



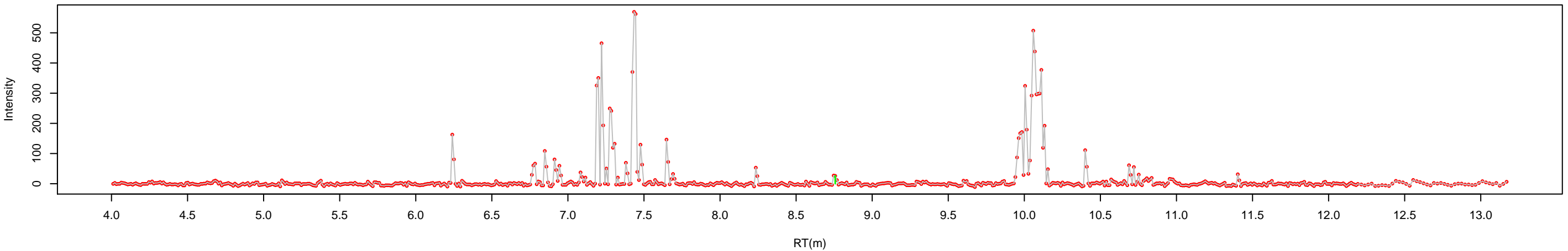
raw chromatogram | batch: 1 sample: std2-10ul-1 conc: 0.005 function: 2 mass: 514.35>79.93



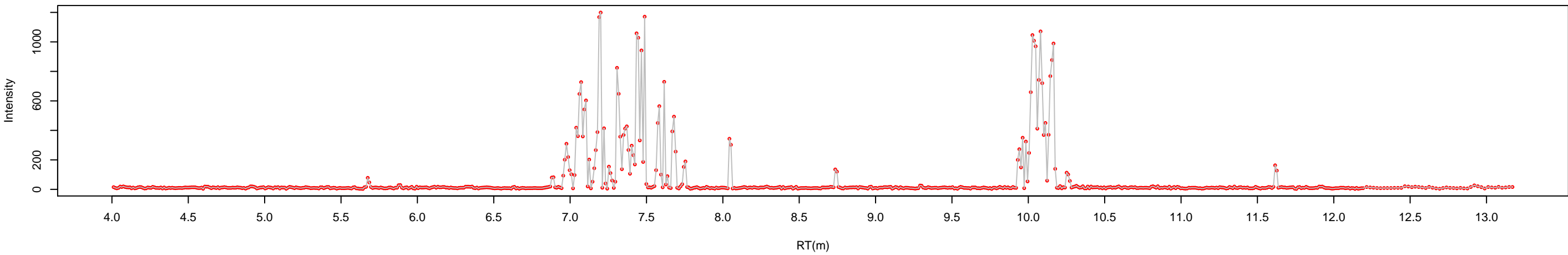
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.0712 BLine: yes



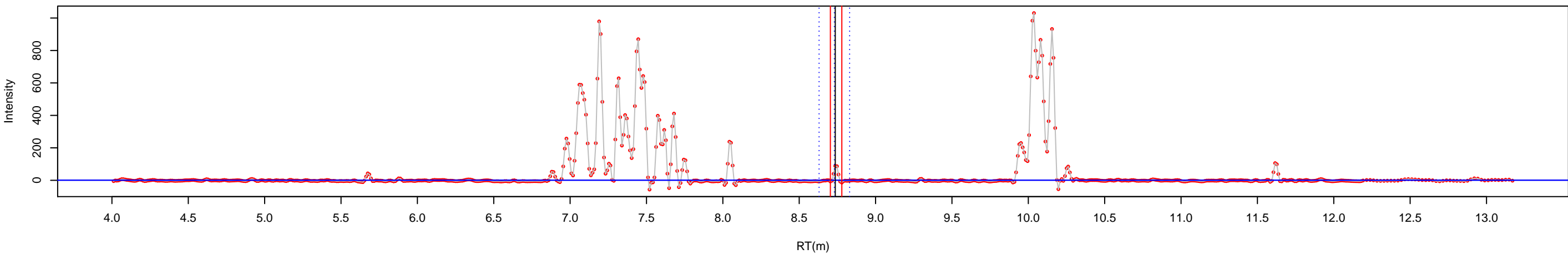
peak area | window size:3 BLine: yes



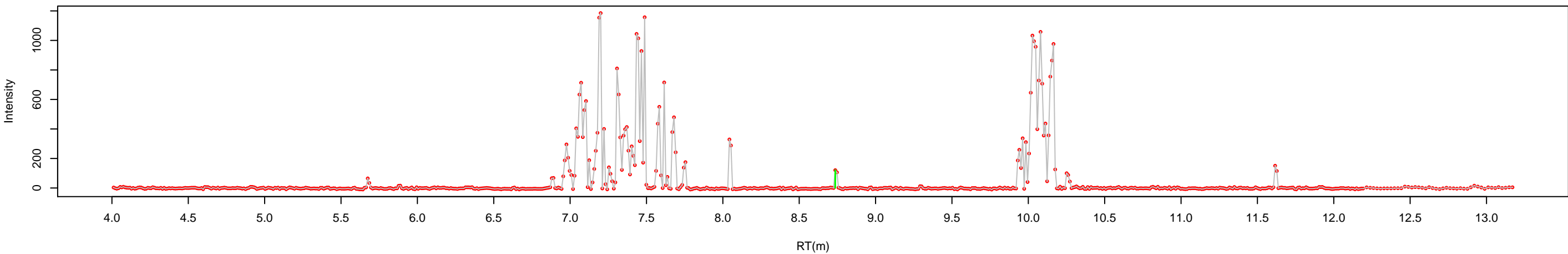
raw chromatogram | batch: 1 sample: std3-10ul-1 conc: 0.01 function: 2 mass: 514.35>79.93



peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.117 BLine: yes

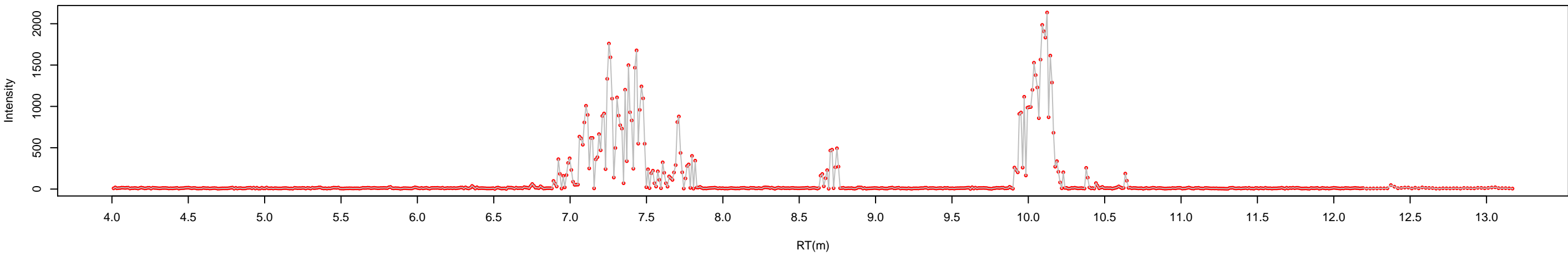


peak area | window size:3 BLine: yes

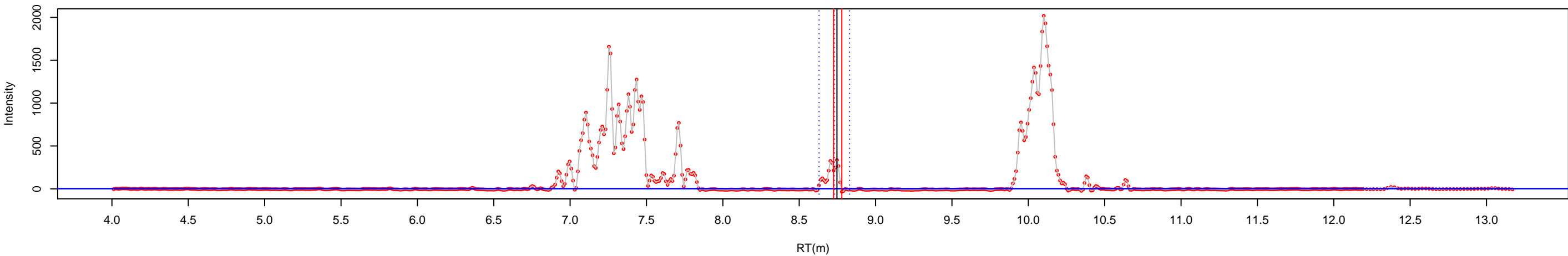




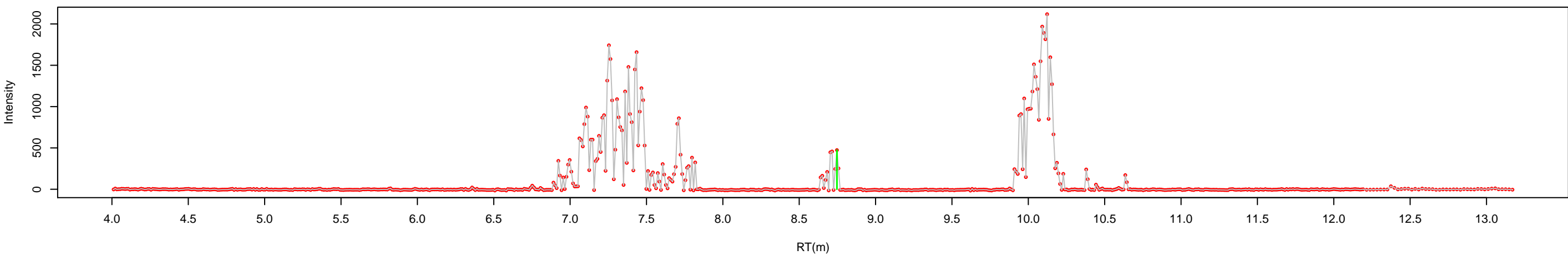
raw chromatogram | batch: 1 sample: std4-10ul-1 conc: 0.02 function: 2 mass: 514.35>79.93



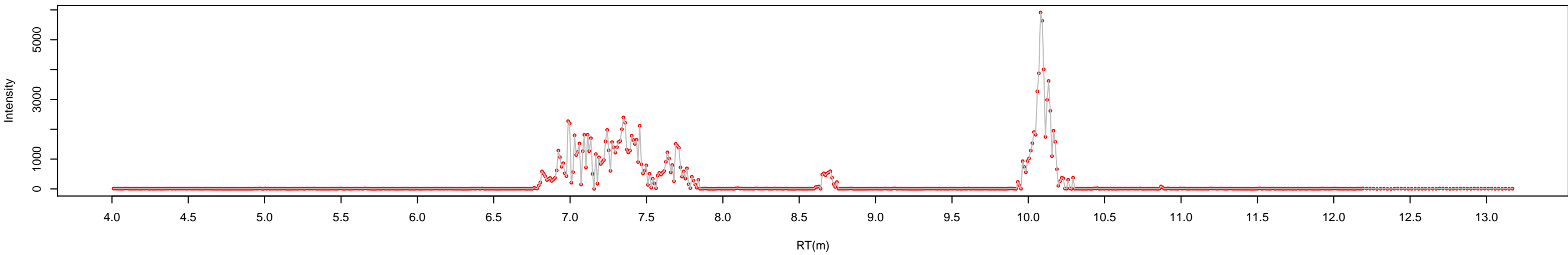
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.41 BLine: yes



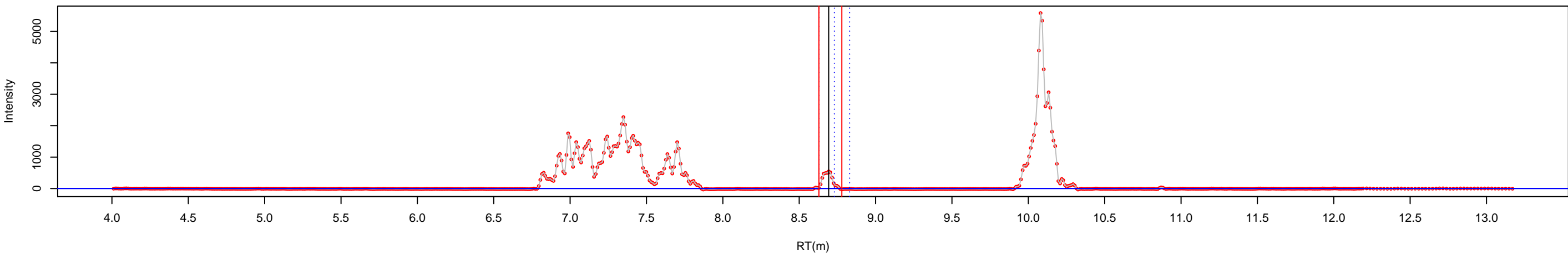
peak area | window size:3 BLine: yes



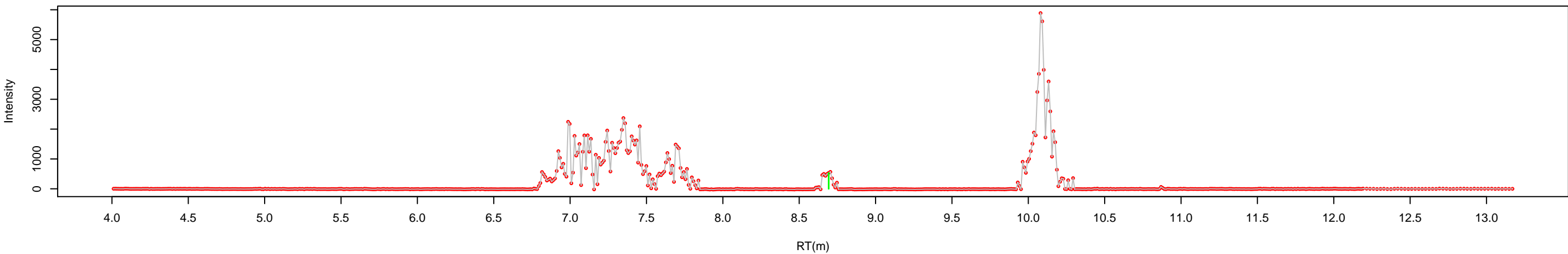
raw chromatogram | batch: 1 sample: std5-10ul-1 conc: 0.05 function: 2 mass: 514.35>79.93



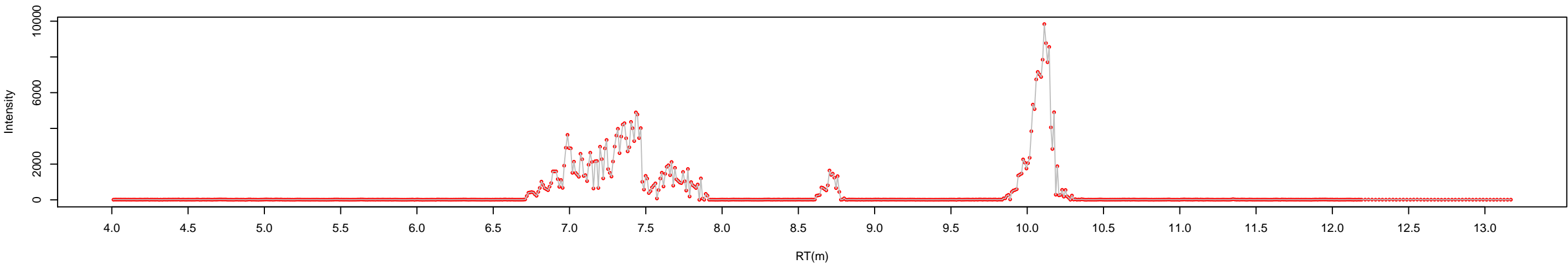
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.271 BLine: yes



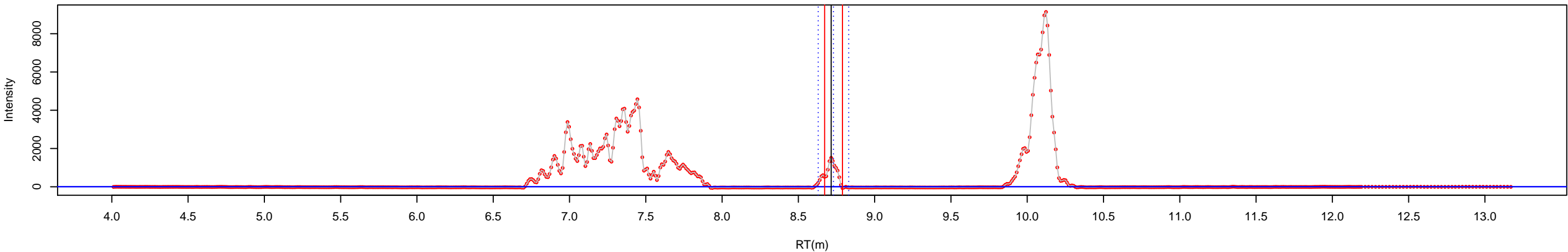
peak area | window size:3 BLine: yes



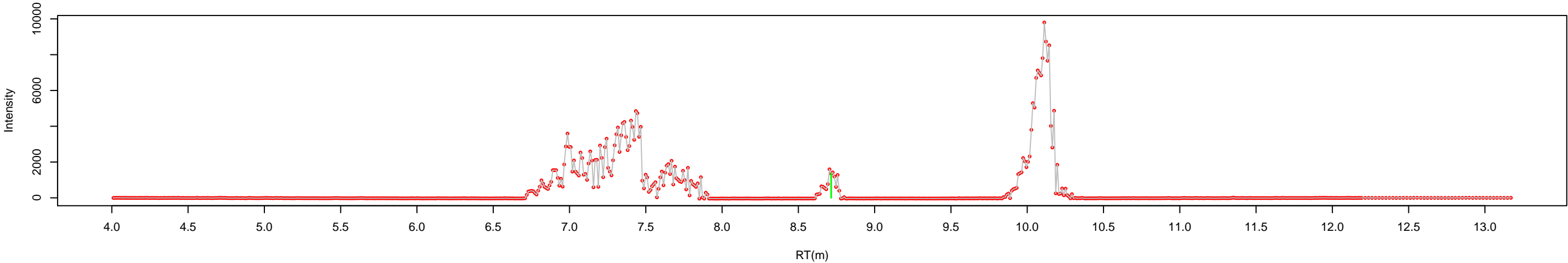
raw chromatogram | batch: 1 sample: std6-10ul-1 conc: 0.1 function: 2 mass: 514.35>79.93



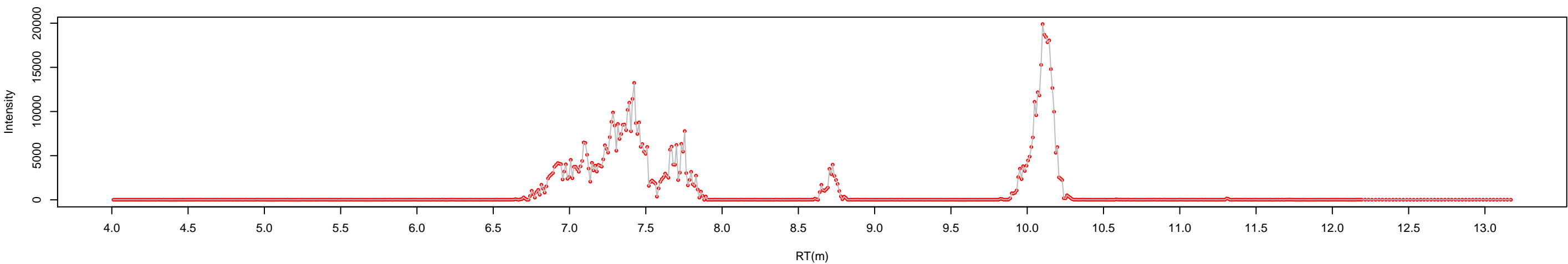
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.79 BLine: yes



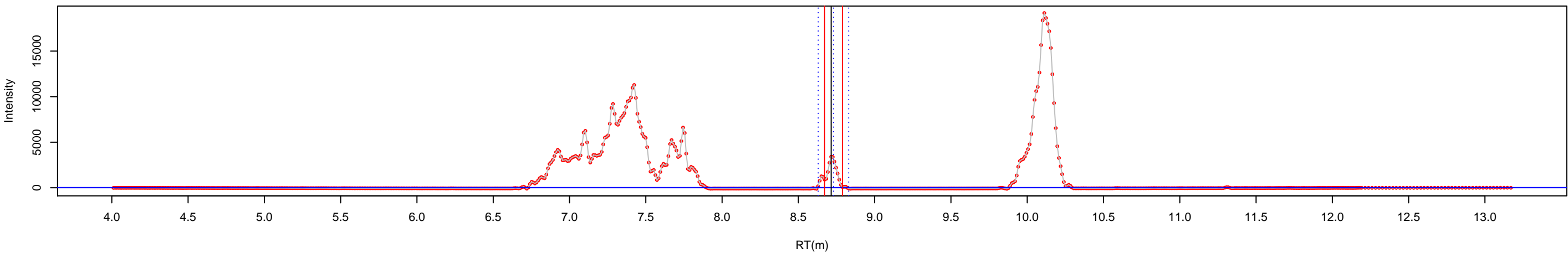
peak area | window size:3 BLine: yes



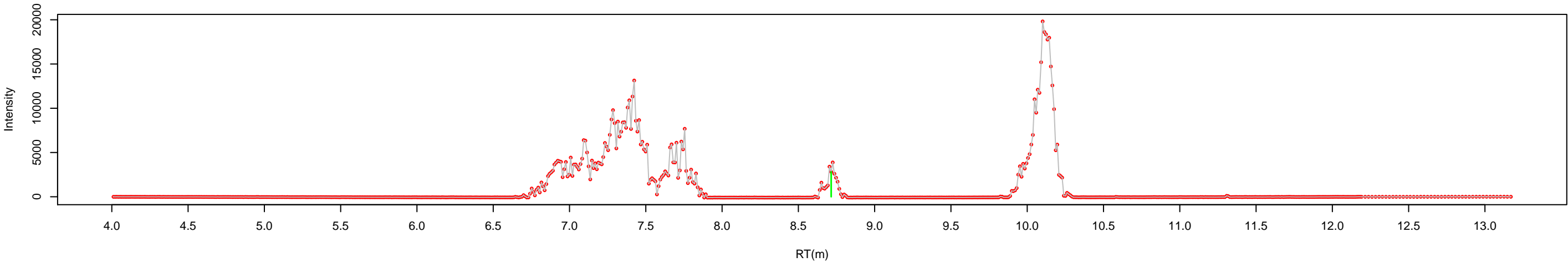
raw chromatogram | batch: 1 sample: std7-10ul-1 conc: 0.2 function: 2 mass: 514.35>79.93



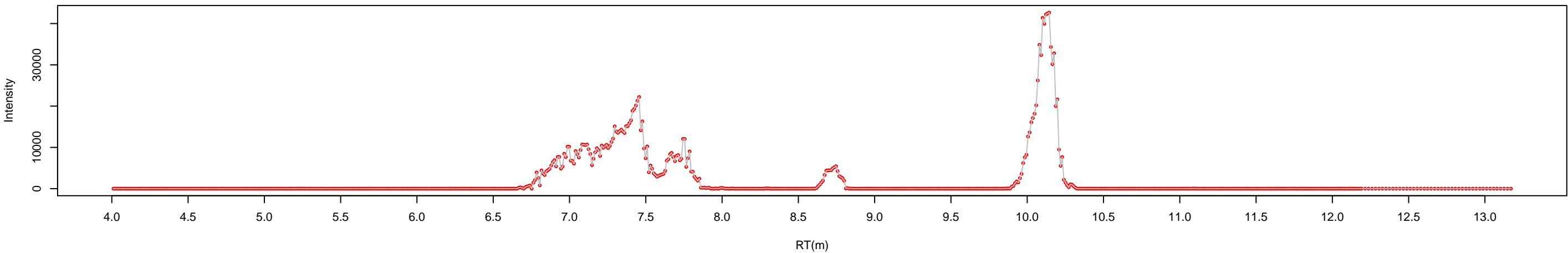
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 2.18 BLine: yes



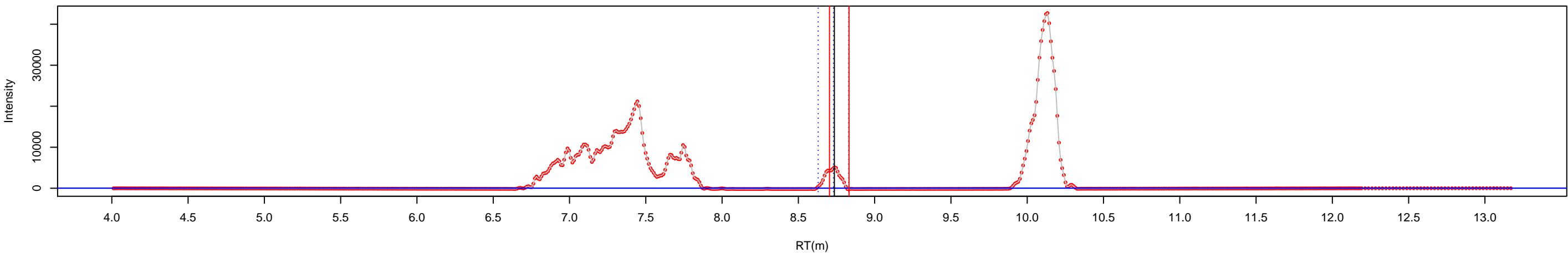
peak area | window size:3 BLine: yes



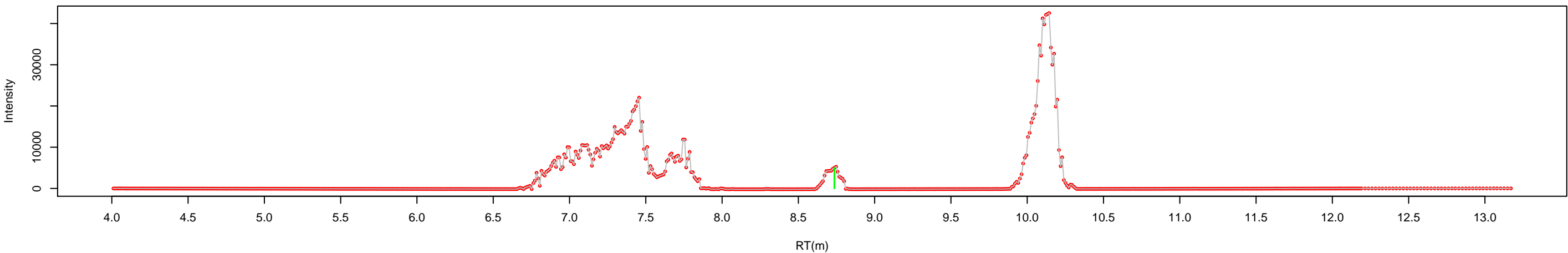
raw chromatogram | batch: 1 sample: std8-10ul-1 conc: 0.5 function: 2 mass: 514.35>79.93



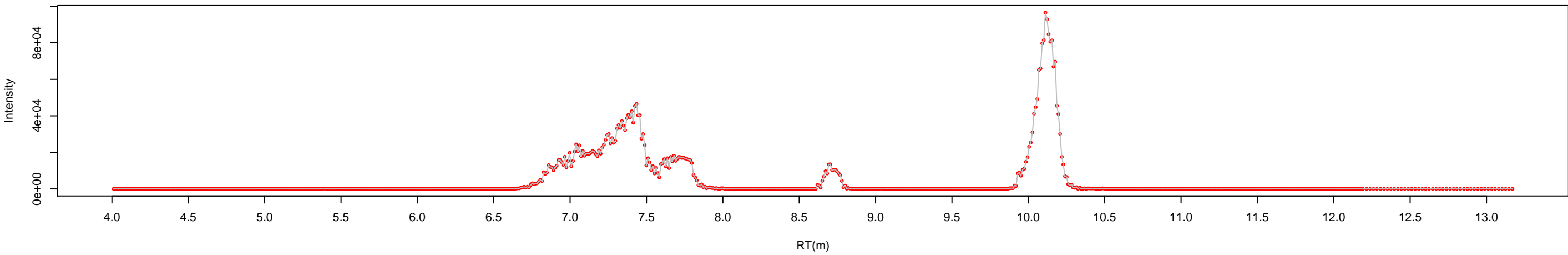
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 4.07 BLine: yes



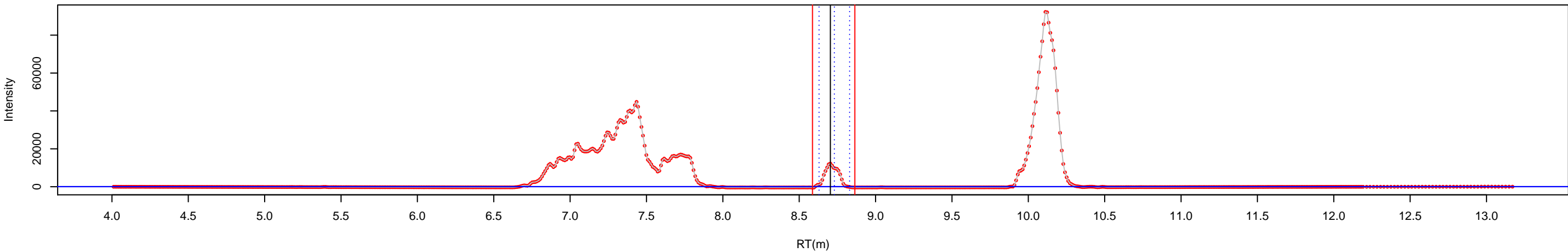
peak area | window size:3 BLine: yes



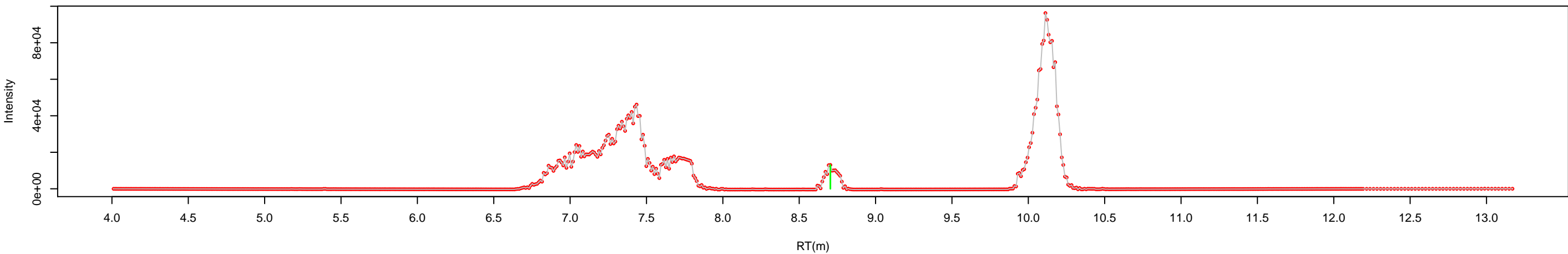
raw chromatogram | batch: 1 sample: std9-10ul-1 conc: 1 function: 2 mass: 514.35>79.93



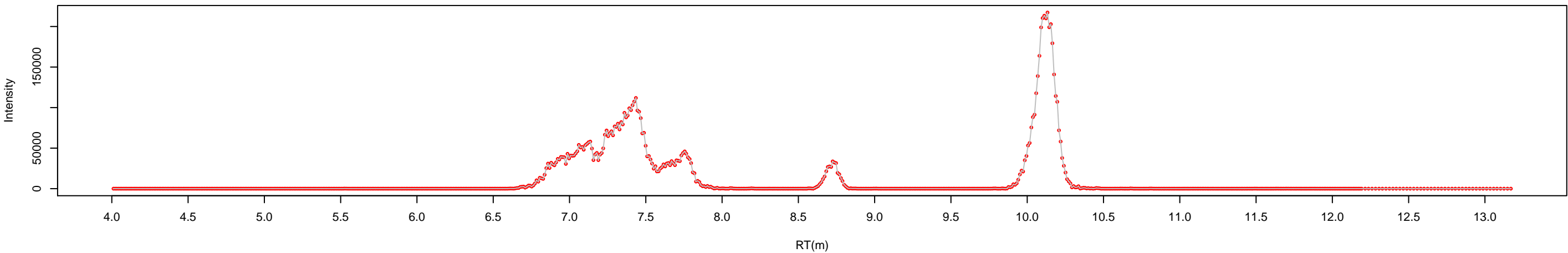
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 8.5 BLine: yes



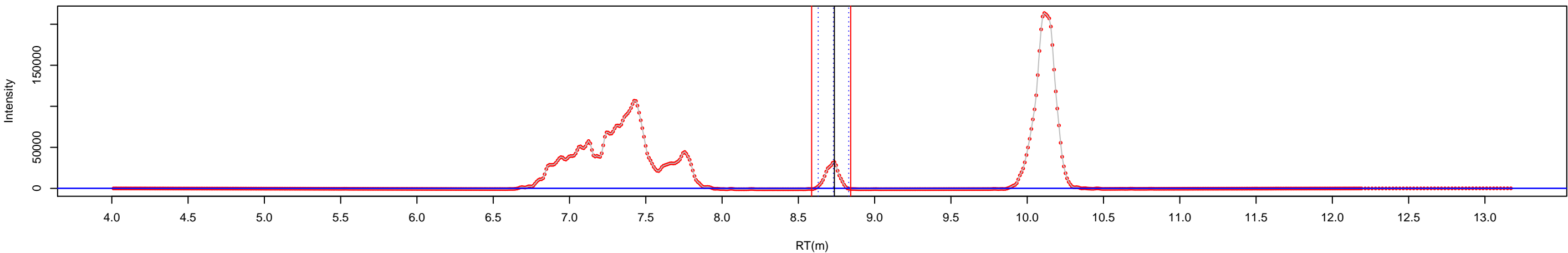
peak area | window size:3 BLine: yes



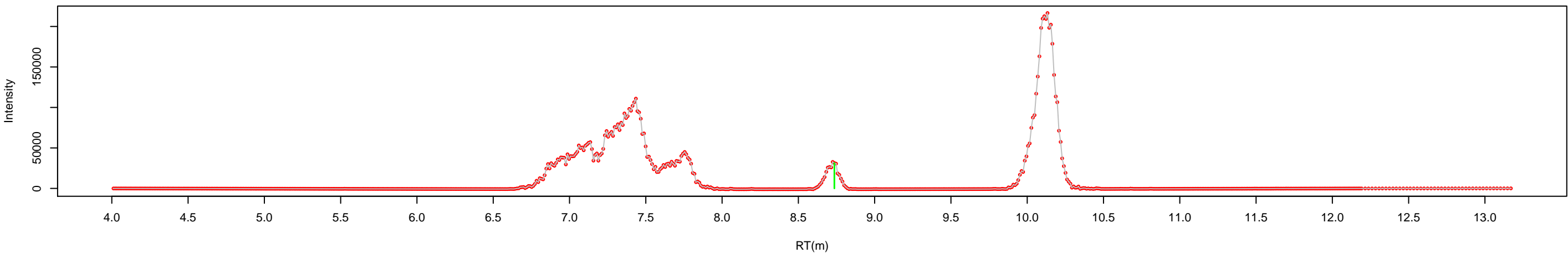
raw chromatogram | batch: 1 sample: std10-10ul-1 conc: 2 function: 2 mass: 514.35>79.93



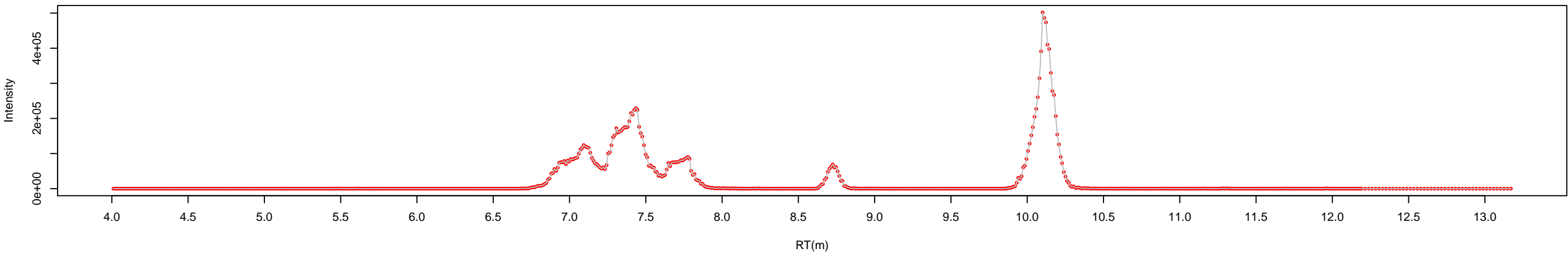
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 19.1 BLine: yes



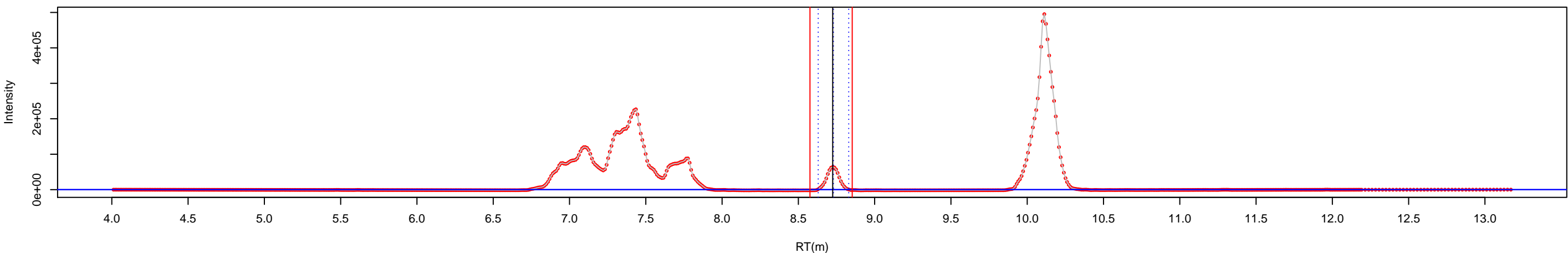
peak area | window size:3 BLine: yes



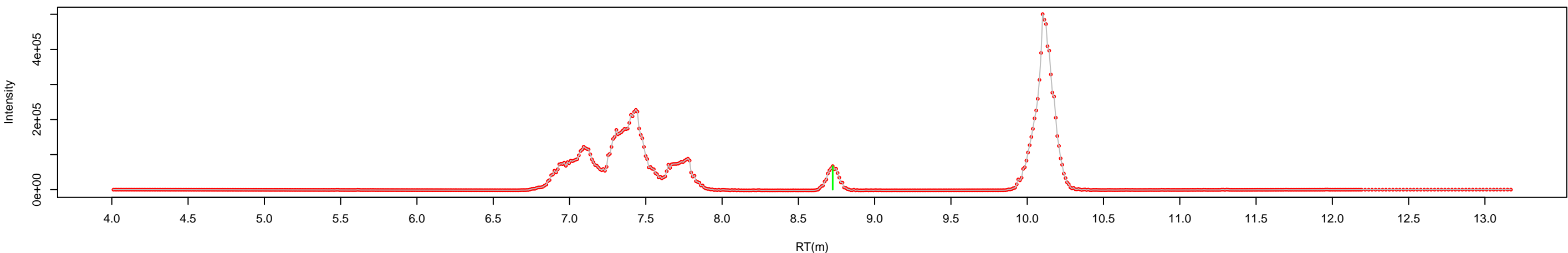
raw chromatogram | batch: 1 sample: std11-10ul-1 conc: 5 function: 2 mass: 514.35>79.93



peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 36.3 BLine: yes

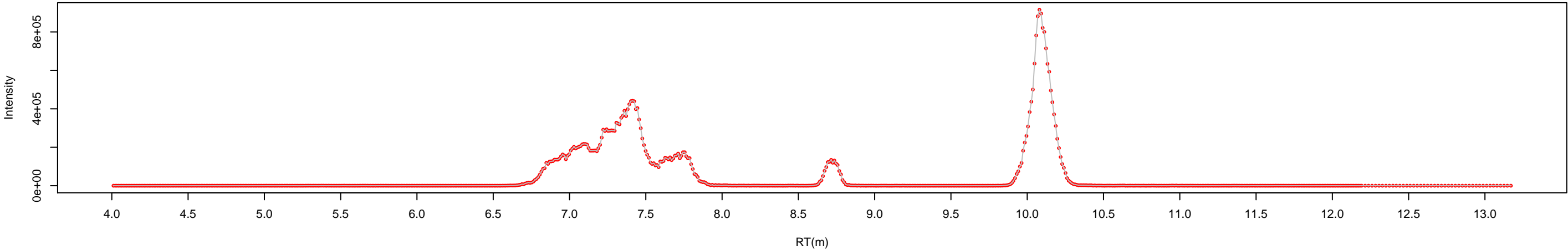


peak area | window size:3 BLine: yes

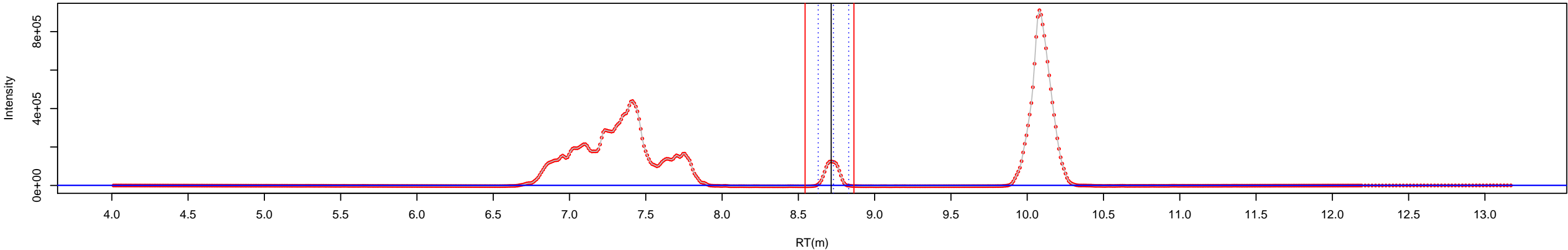




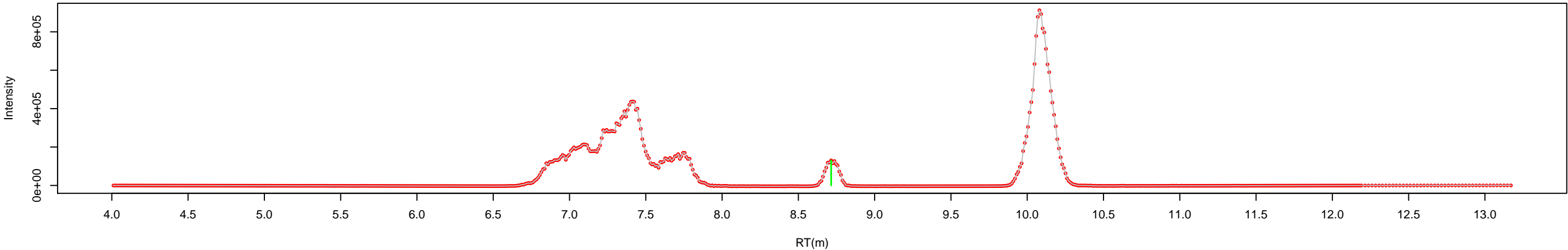
raw chromatogram | batch: 1   sample: std12-10ul-1   conc: 10   function: 2   mass: 514.35>79.93



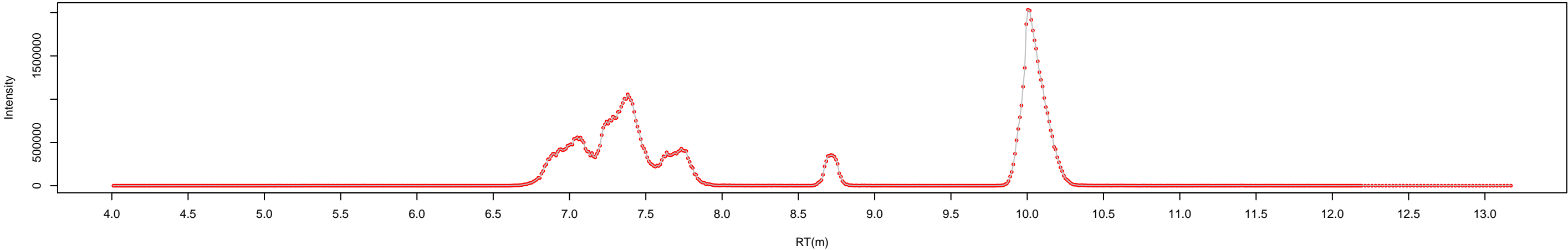
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 98.4   BLine: yes



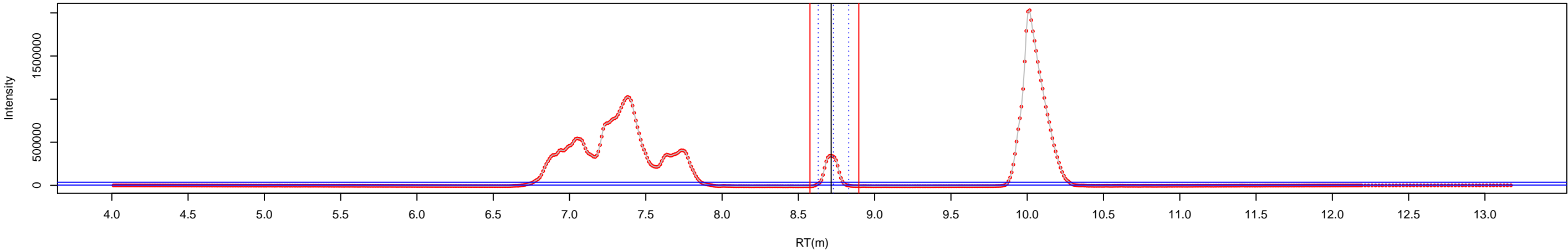
peak area | window size:3   BLine: yes



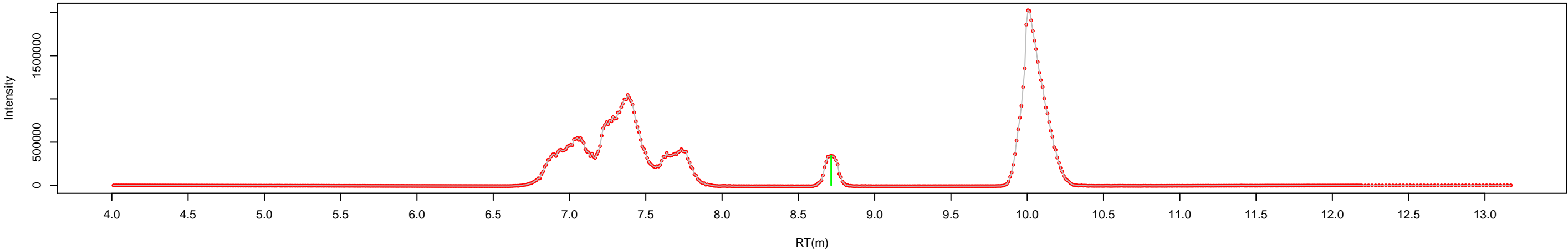
raw chromatogram | batch: 1   sample: std13-10ul-1   conc: 20   function: 2   mass: 514.35>79.93



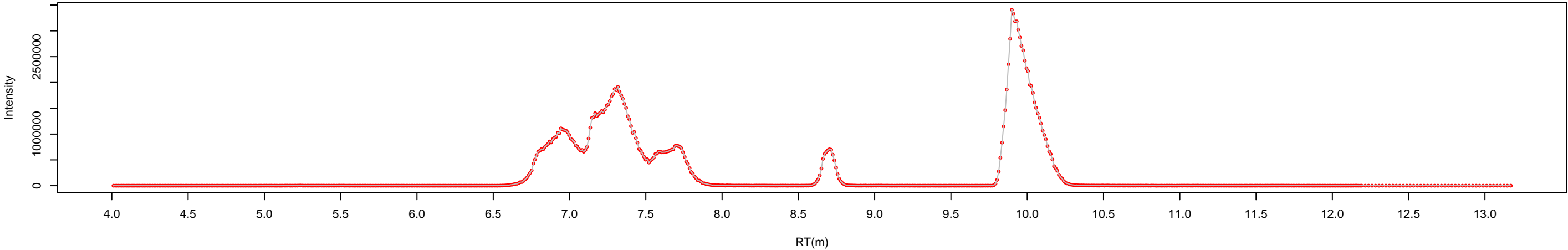
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 3630   BLine: yes



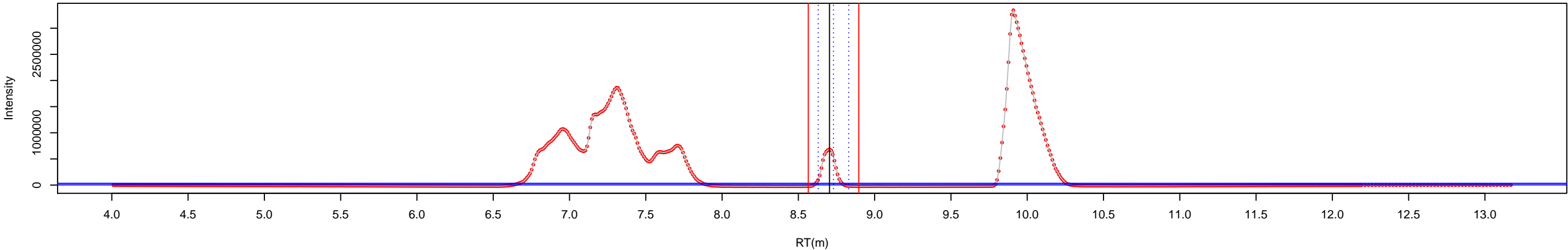
peak area | window size:3   BLine: yes



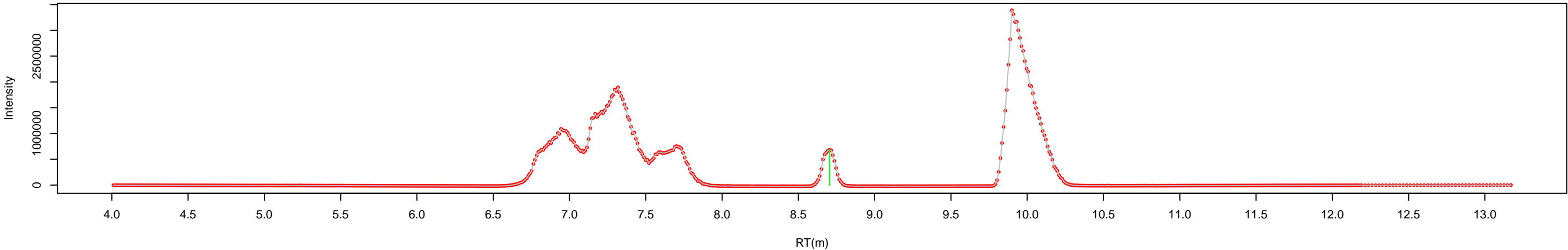
raw chromatogram | batch: 1 sample: std14-10ul-1 conc: 50 function: 2 mass: 514.35>79.93



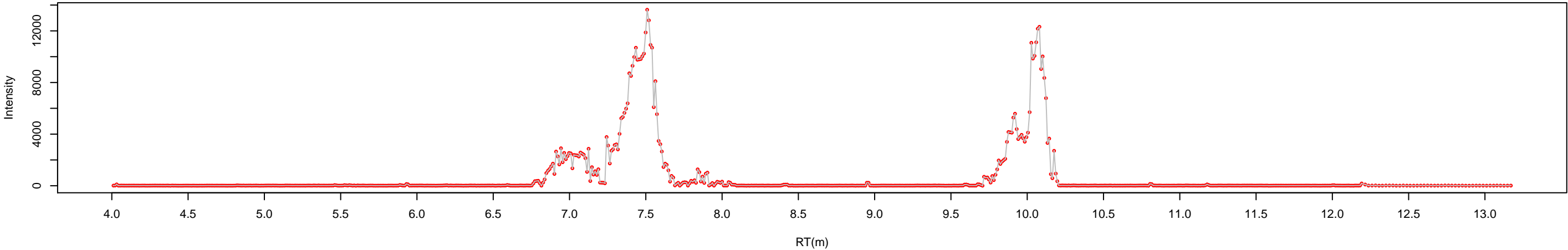
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 3290 BLine: yes



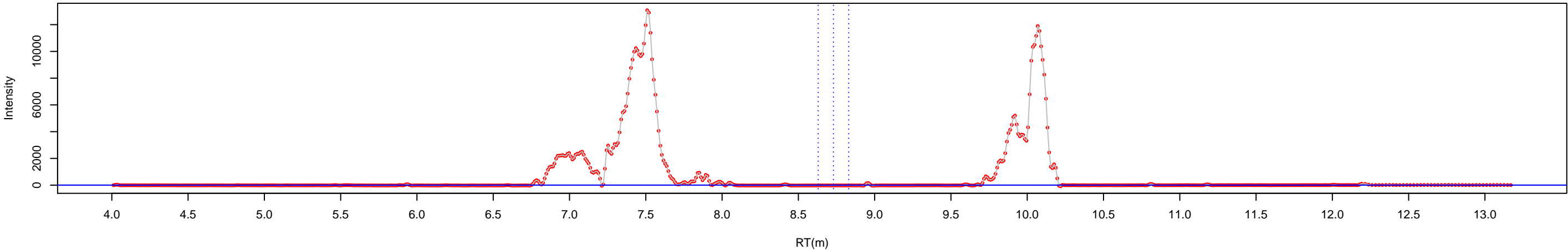
peak area | window size:3 BLine: yes



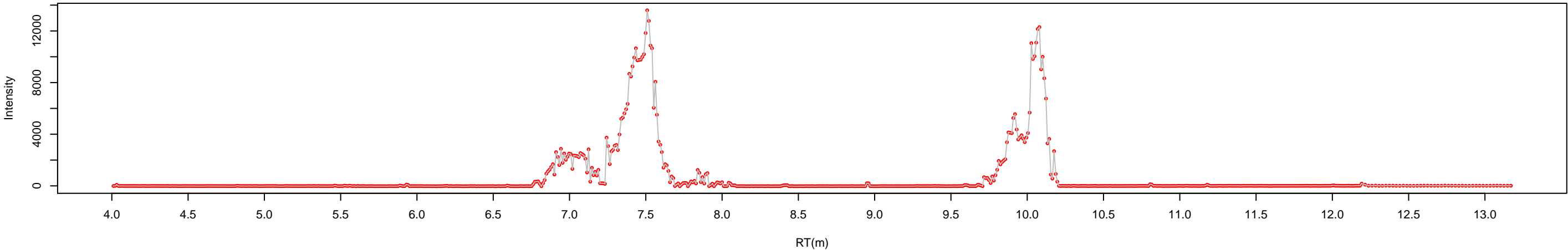
raw chromatogram | batch: 2 sample: coth45 conc: NA function: 2 mass: 514.35>79.93



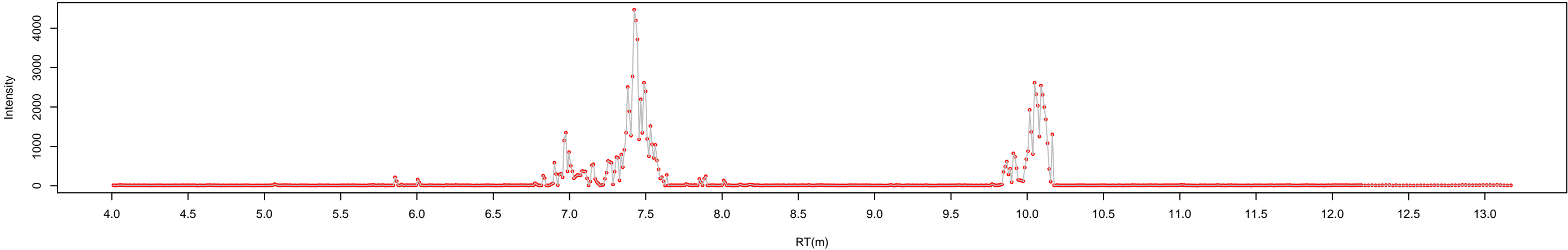
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.609 BLine: yes



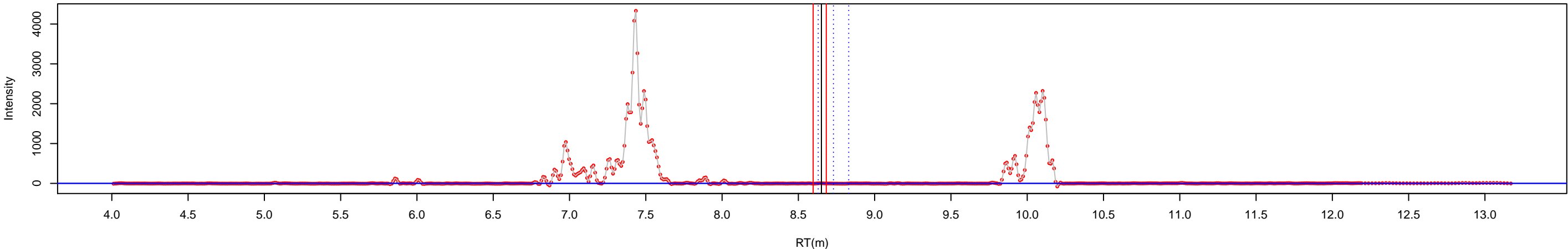
peak area | window size:3 BLine: yes



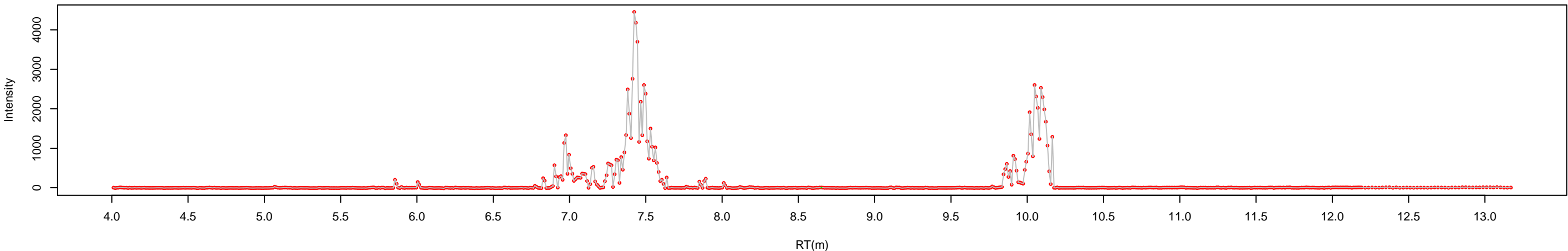
raw chromatogram | batch: 2 sample: coth46 conc: NA function: 2 mass: 514.35>79.93



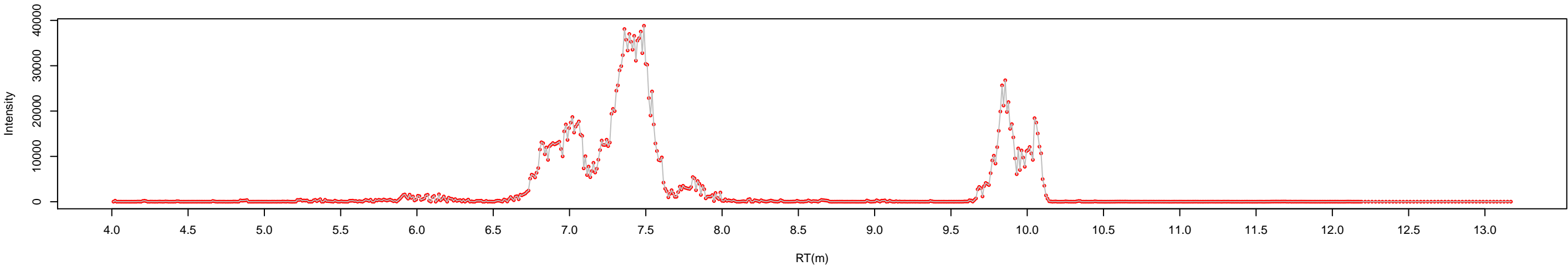
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.0967 BLine: yes



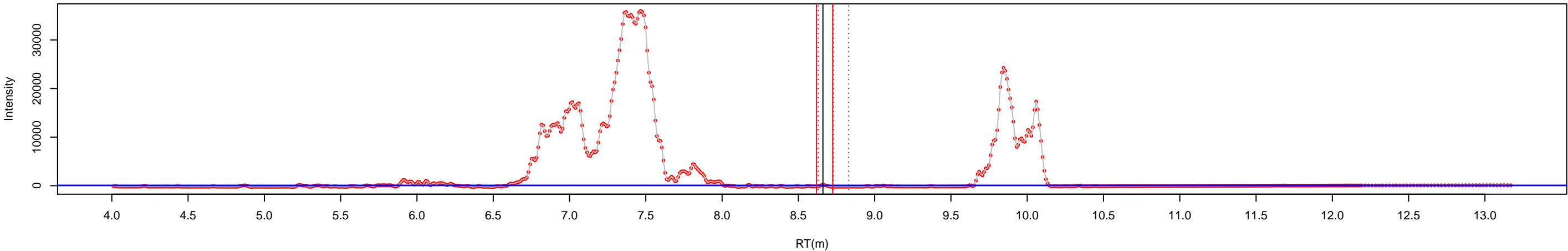
peak area | window size:3 BLine: yes



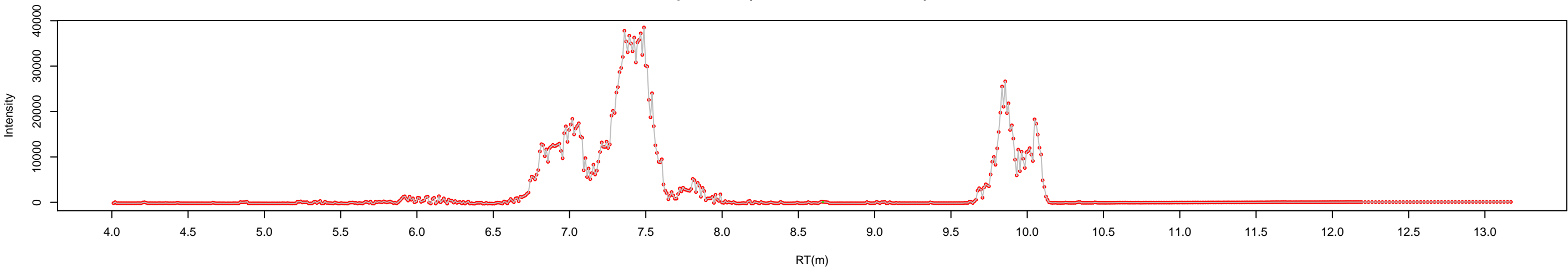
raw chromatogram | batch: 2 sample: coth47 conc: NA function: 2 mass: 514.35>79.93



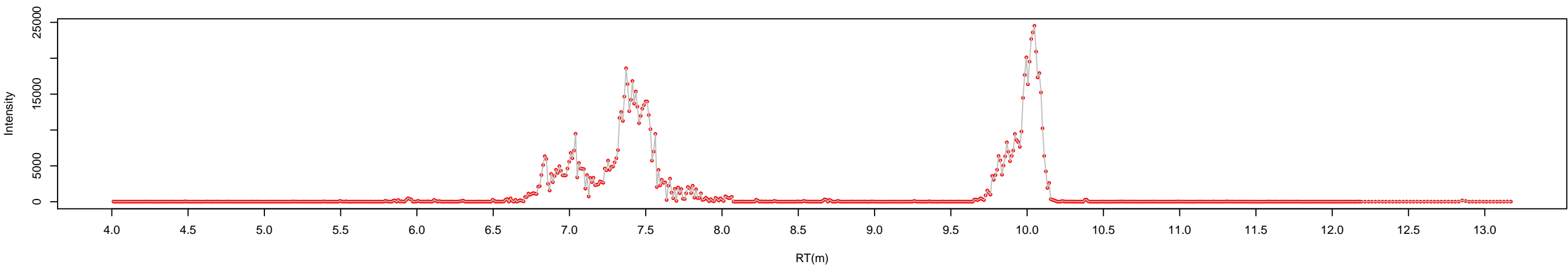
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 10.1 BLine: yes



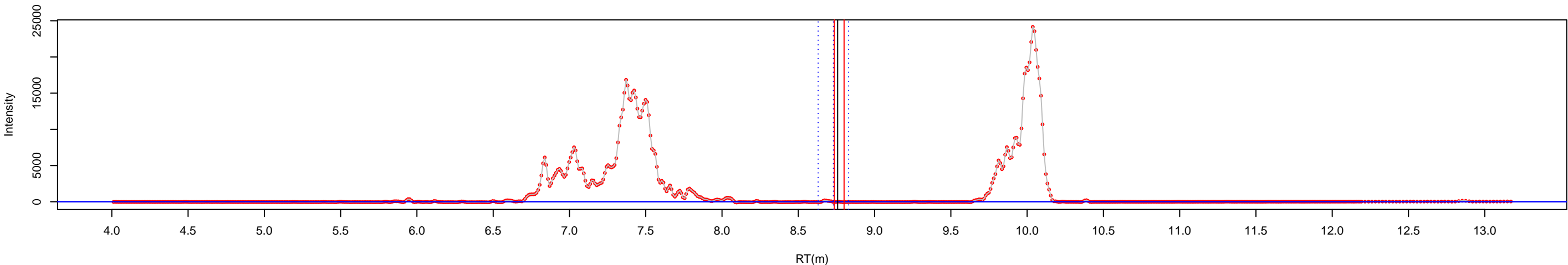
peak area | window size:3 BLine: yes



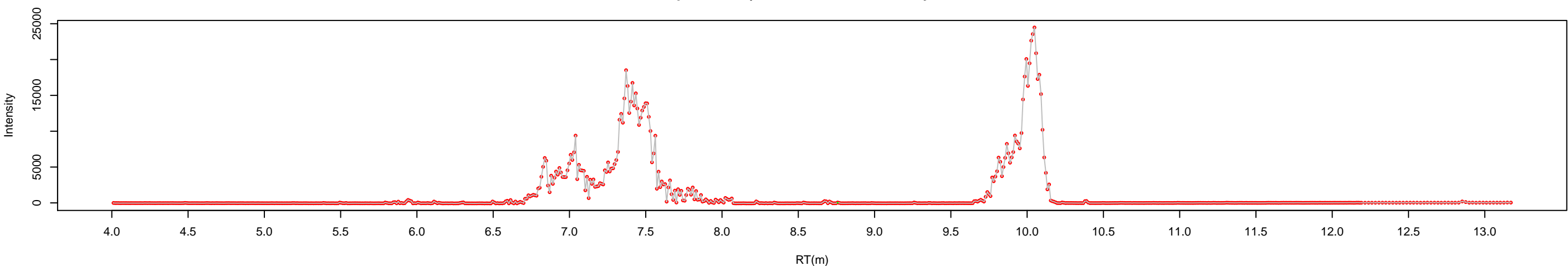
raw chromatogram | batch: 2 sample: coth48 conc: NA function: 2 mass: 514.35>79.93



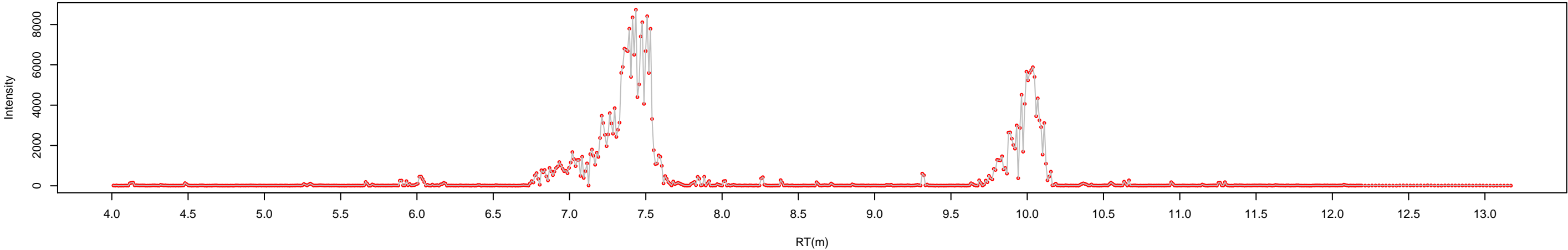
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 1.09 BLine: yes



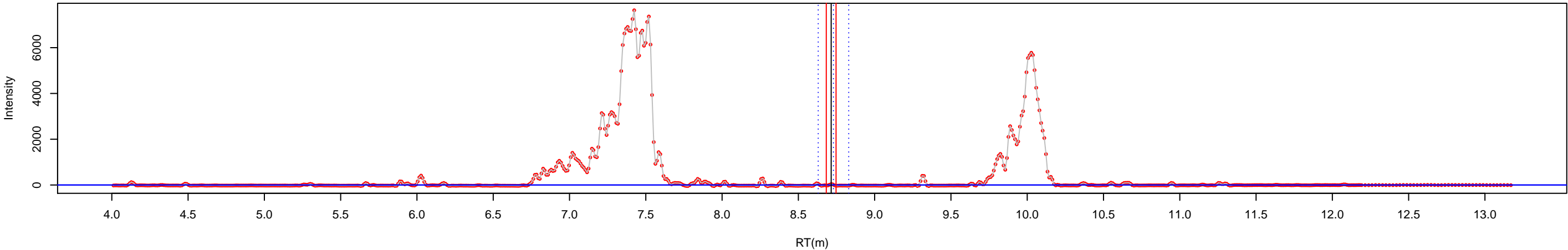
peak area | window size:3 BLine: yes



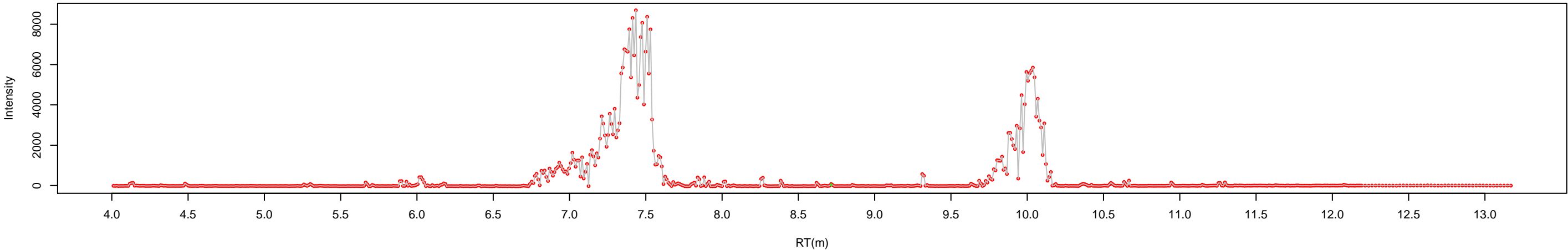
raw chromatogram | batch: 2 sample: coth49 conc: NA function: 2 mass: 514.35>79.93



peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.224 BLine: yes

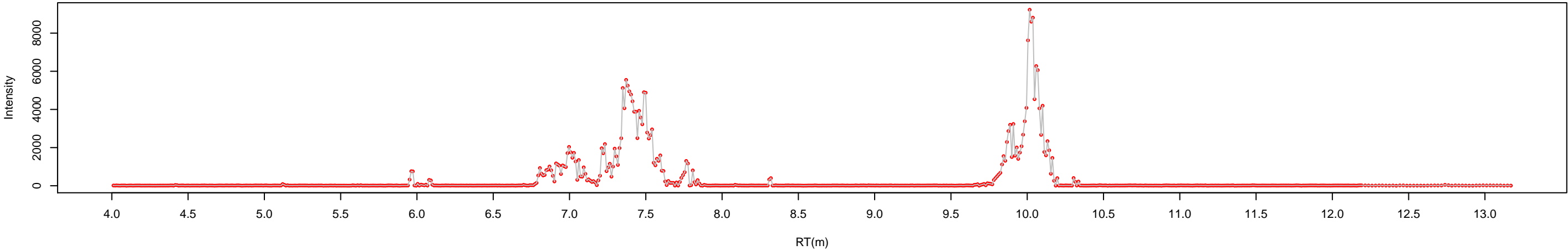


peak area | window size:3 BLine: yes

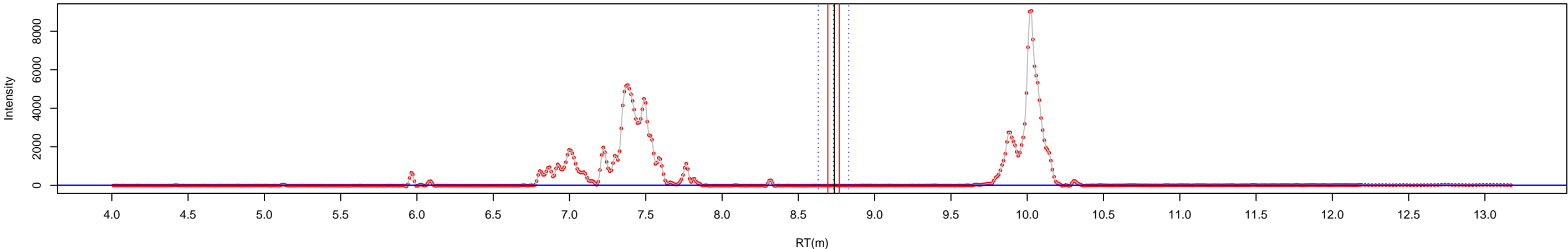




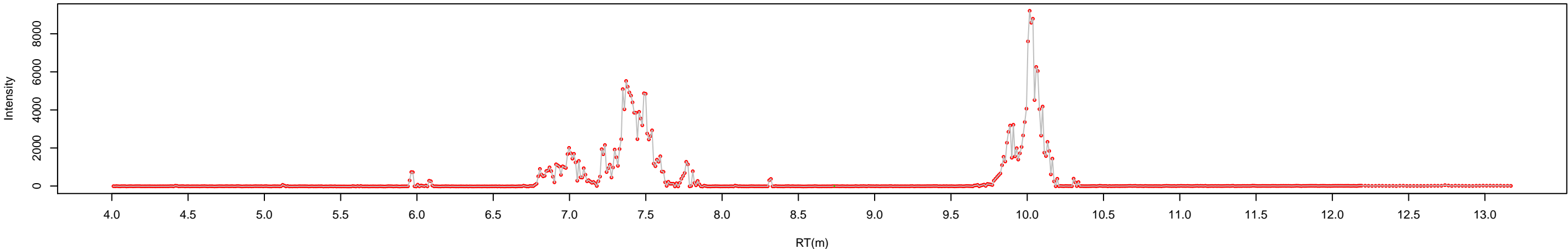
raw chromatogram | batch: 2 sample: coth50 conc: NA function: 2 mass: 514.35>79.93



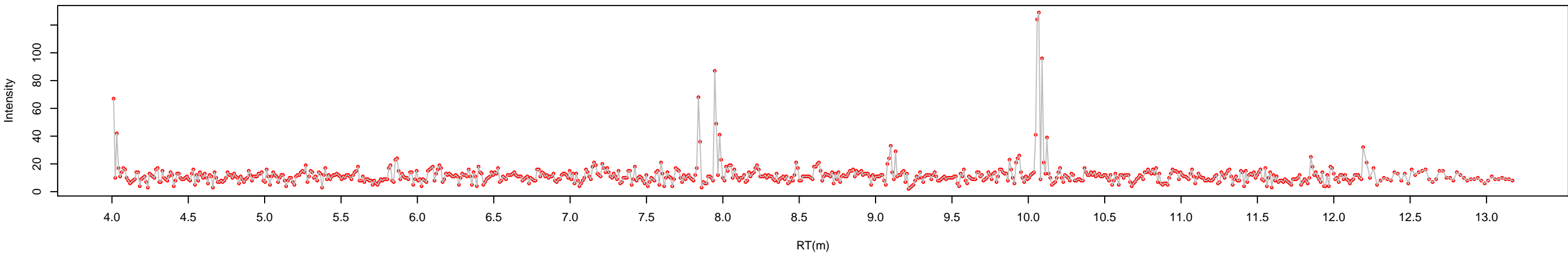
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.336 BLine: yes



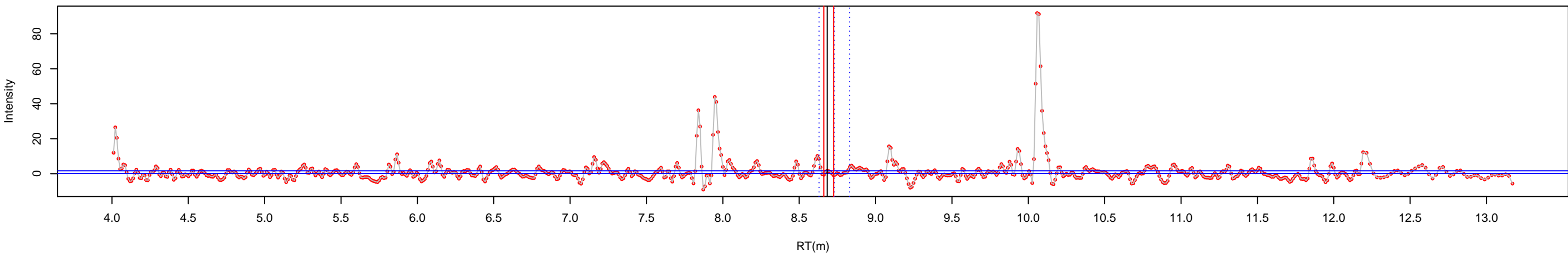
peak area | window size:3 BLine: yes



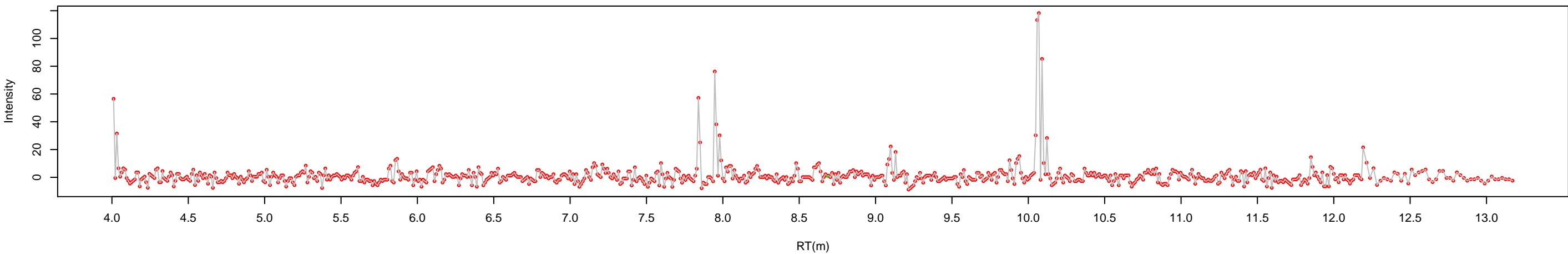
raw chromatogram | batch: 2 sample: std0-10ul-2 conc: 0.001 function: 2 mass: 514.35>79.93



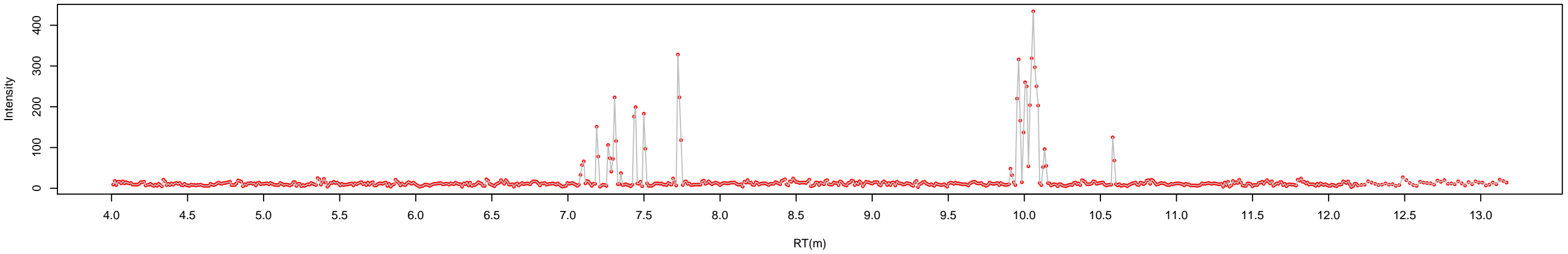
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.164 BLine: yes



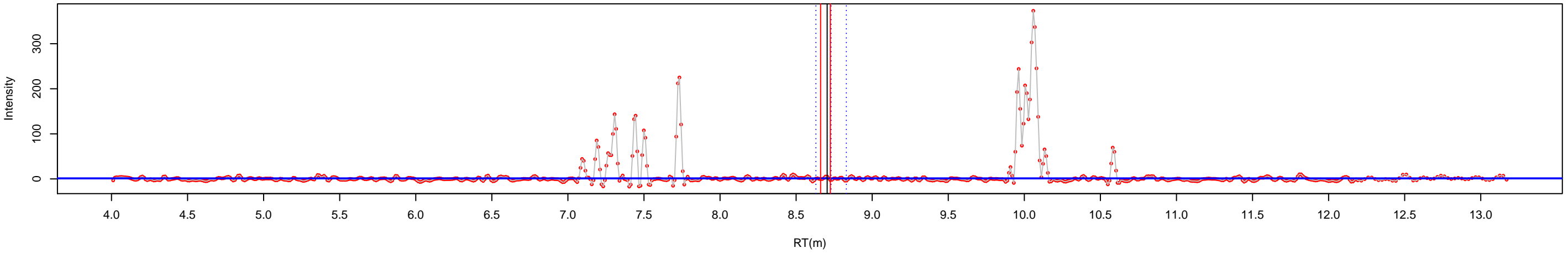
peak area | window size:3 BLine: yes



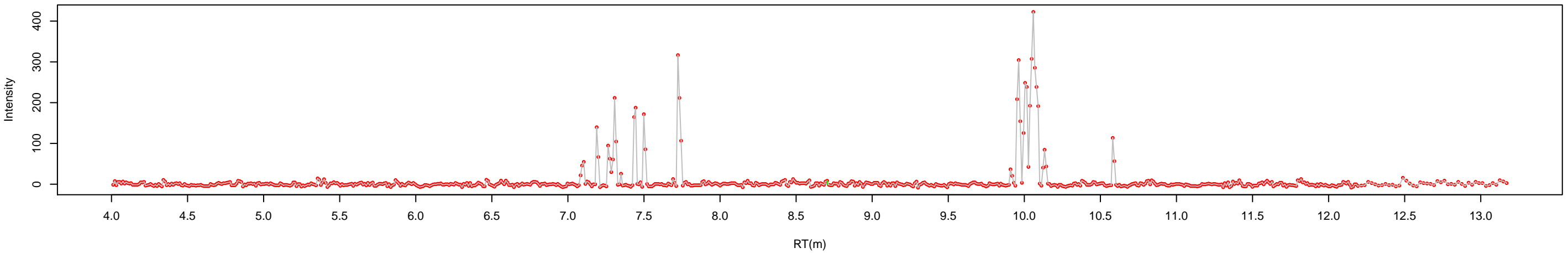
raw chromatogram | batch: 2   sample: std1-10ul-2   conc: 0.002   function: 2   mass: 514.35>79.93



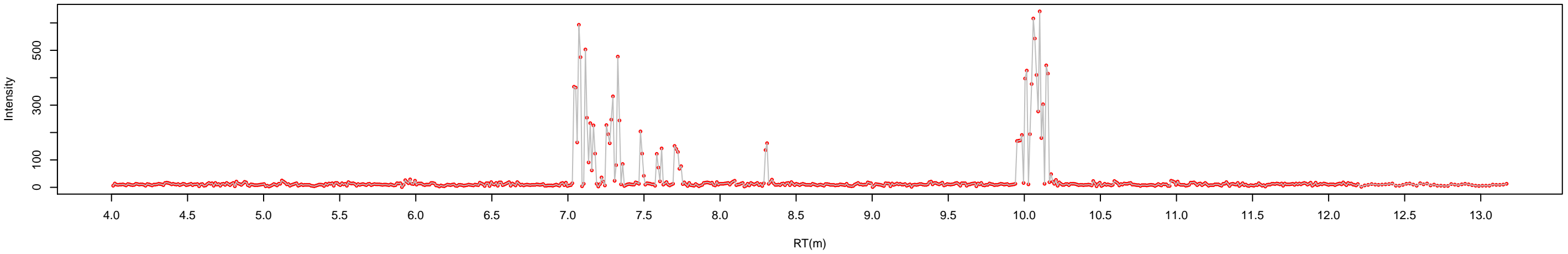
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 0.224   BLine: yes



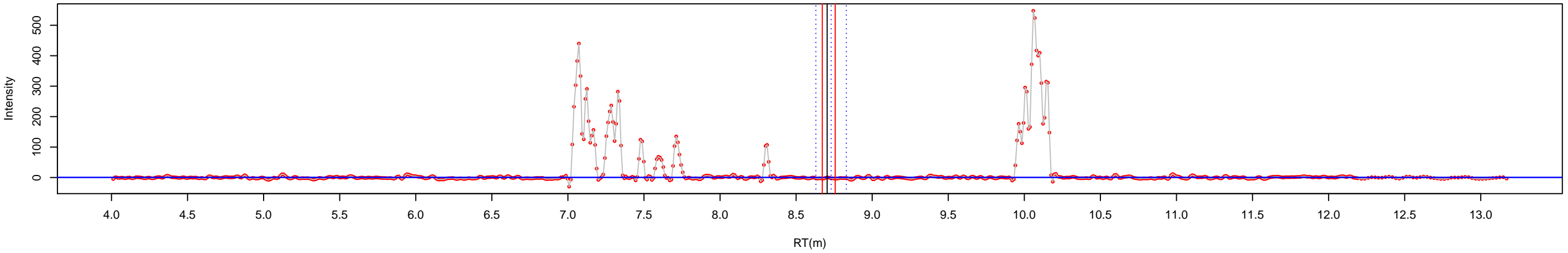
peak area | window size:3   BLine: yes



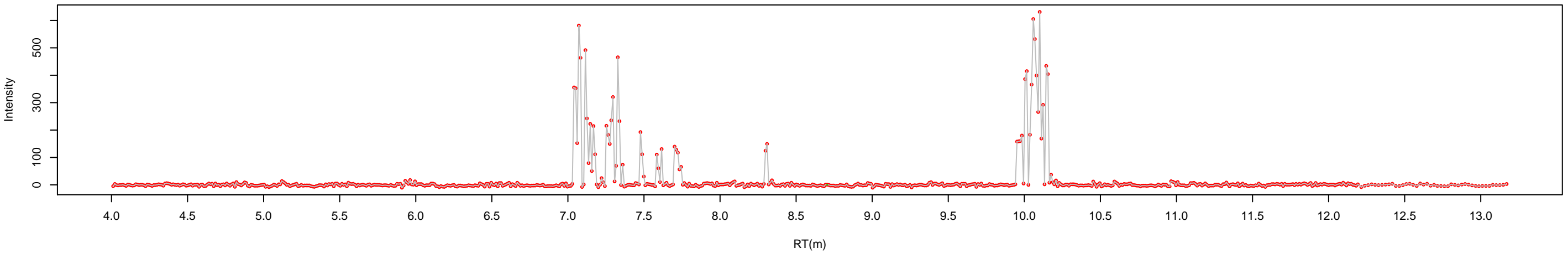
raw chromatogram | batch: 2   sample: std2-10ul-2   conc: 0.005   function: 2   mass: 514.35>79.93



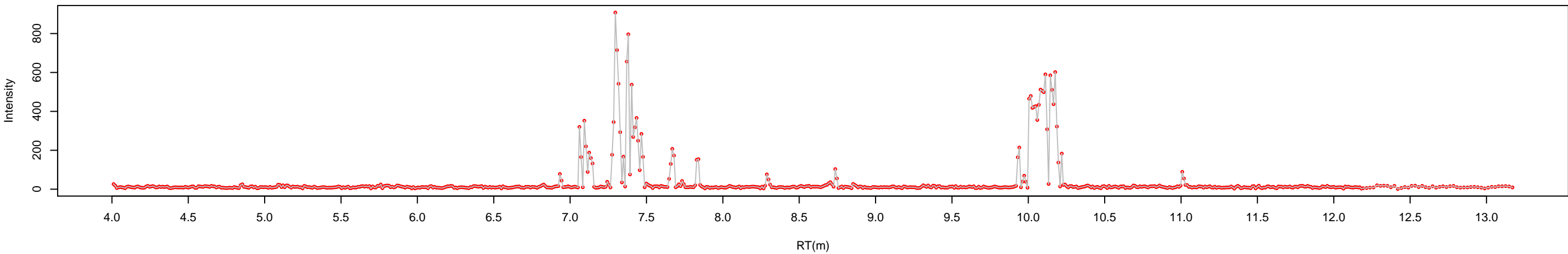
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 0.0741   BLine: yes



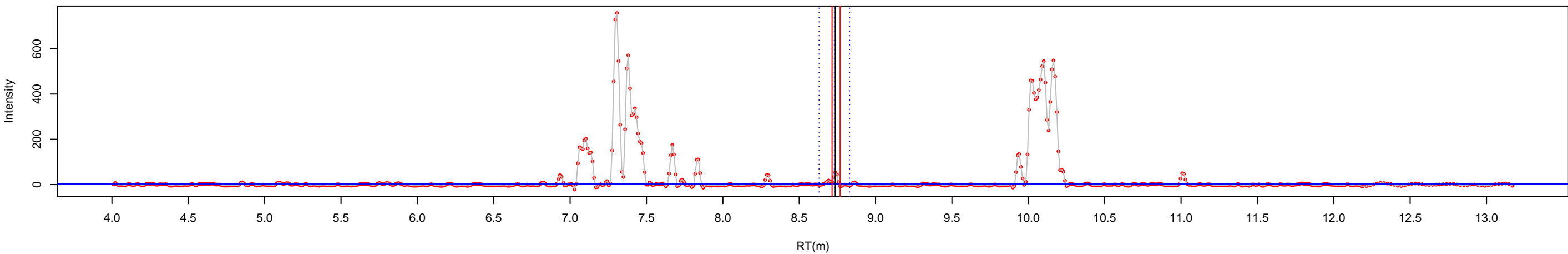
peak area | window size:3   BLine: yes



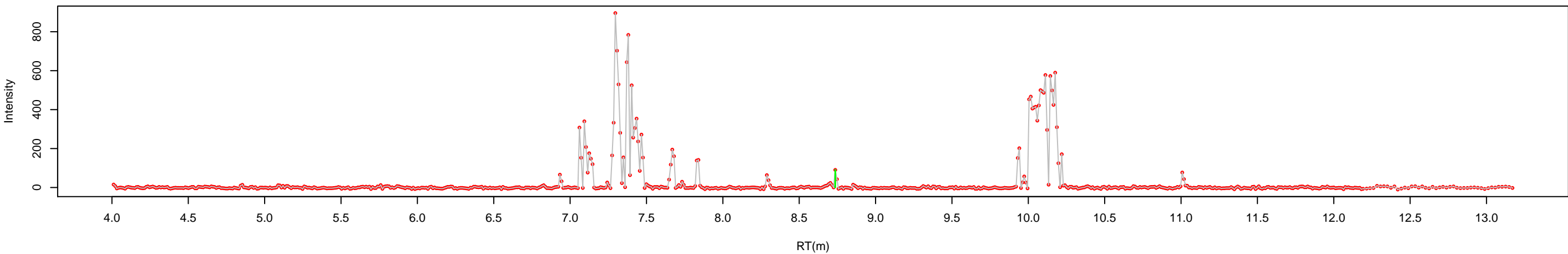
raw chromatogram | batch: 2 sample: std3-10ul-2 conc: 0.01 function: 2 mass: 514.35>79.93



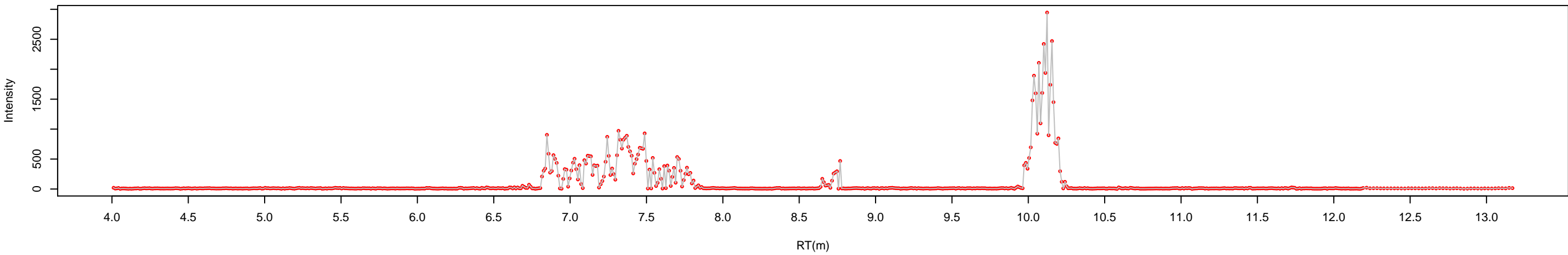
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.295 BLine: yes



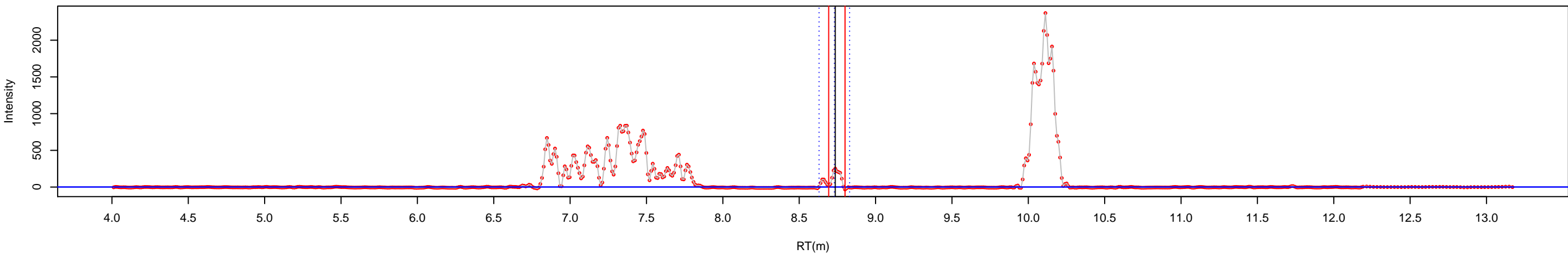
peak area | window size:3 BLine: yes



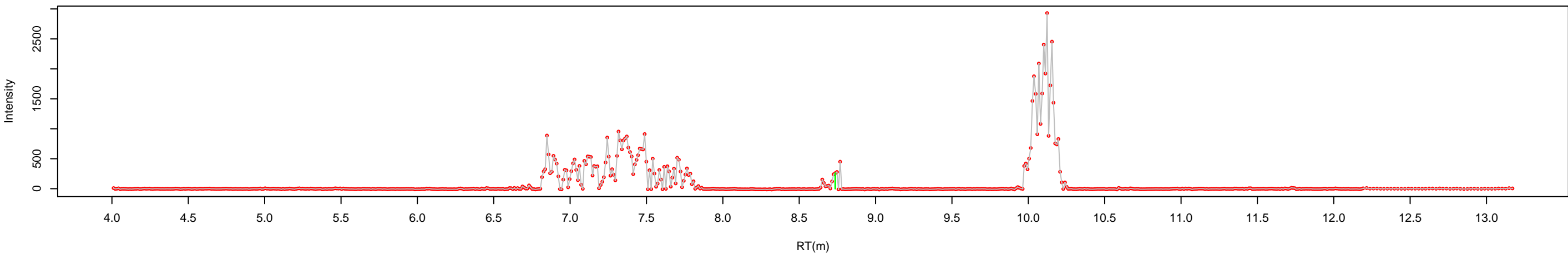
raw chromatogram | batch: 2 sample: std4-10ul-2 conc: 0.02 function: 2 mass: 514.35>79.93



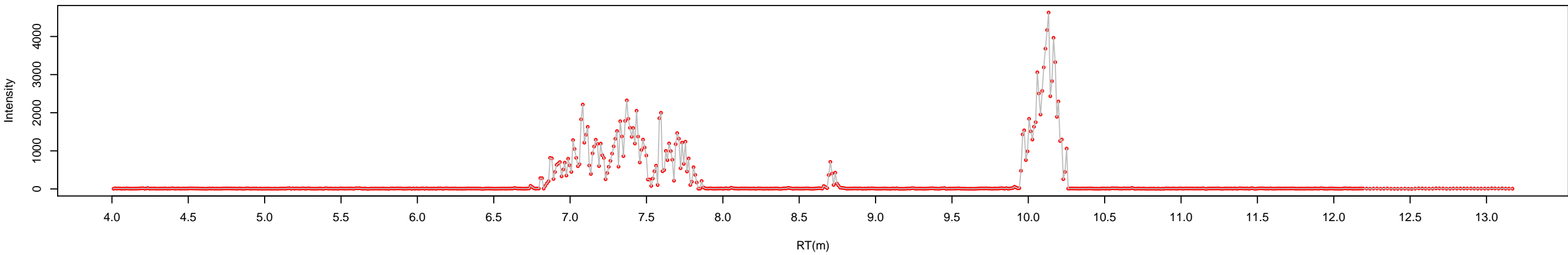
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.158 BLine: yes



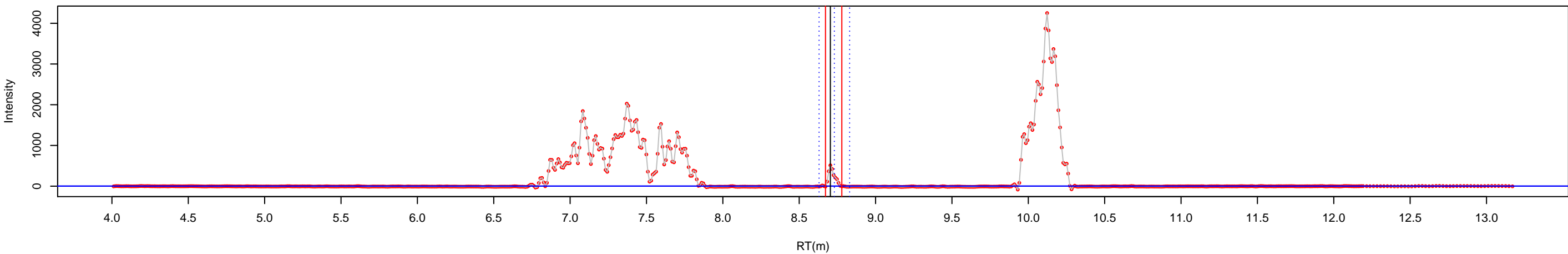
peak area | window size:3 BLine: yes



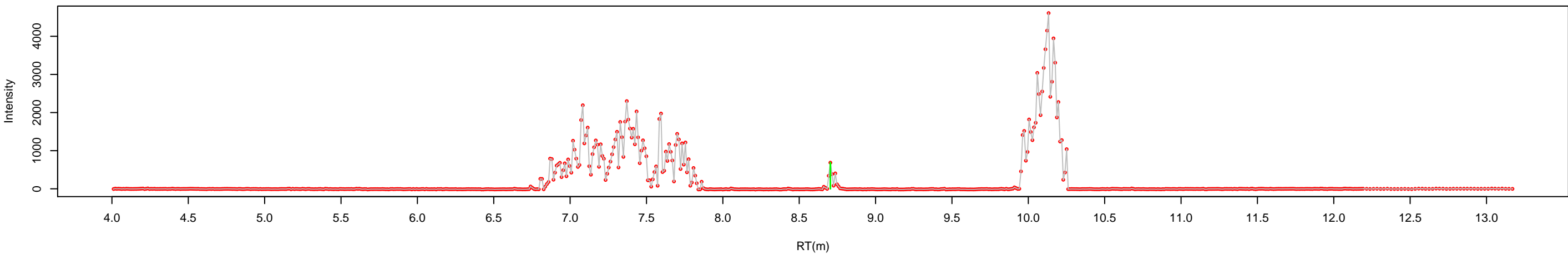
raw chromatogram | batch: 2 sample: std5-10ul-2 conc: 0.05 function: 2 mass: 514.35>79.93



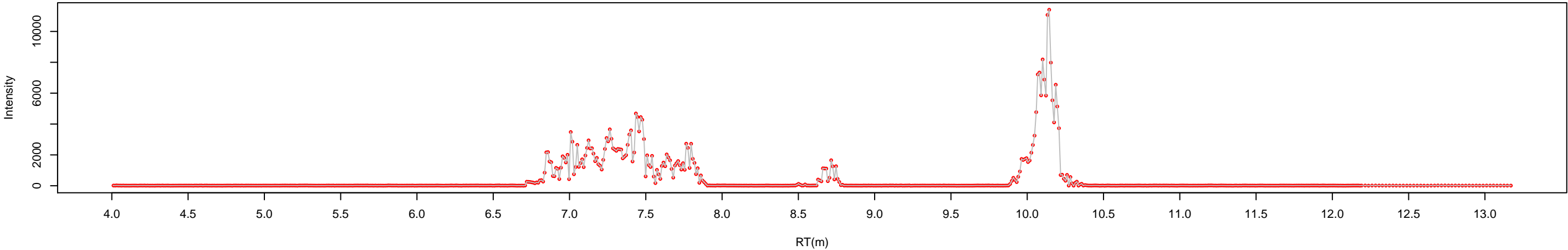
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.378 BLine: yes



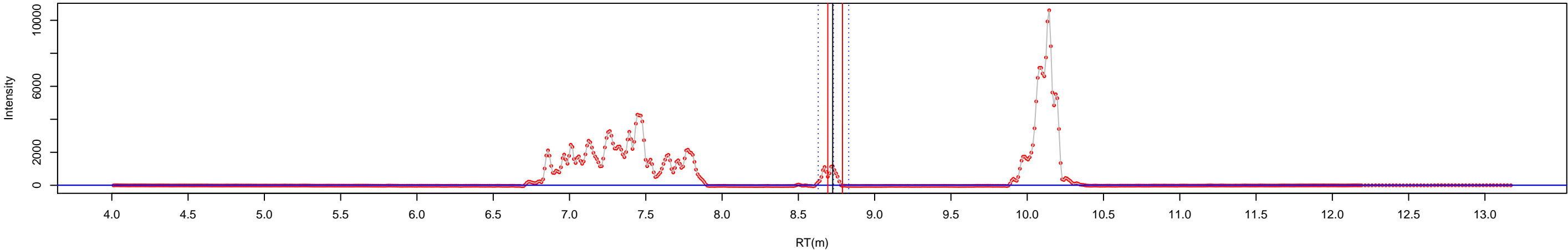
peak area | window size:3 BLine: yes



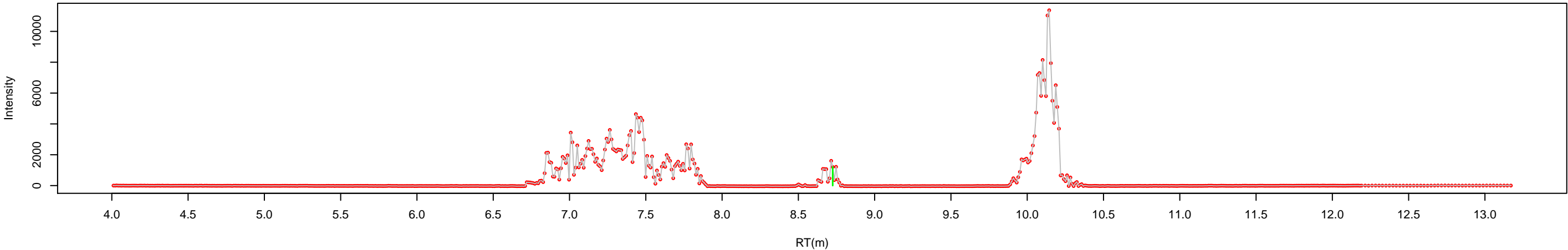
raw chromatogram | batch: 2   sample: std6-10ul-2   conc: 0.1   function: 2   mass: 514.35>79.93



peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 1.09   BLine: yes

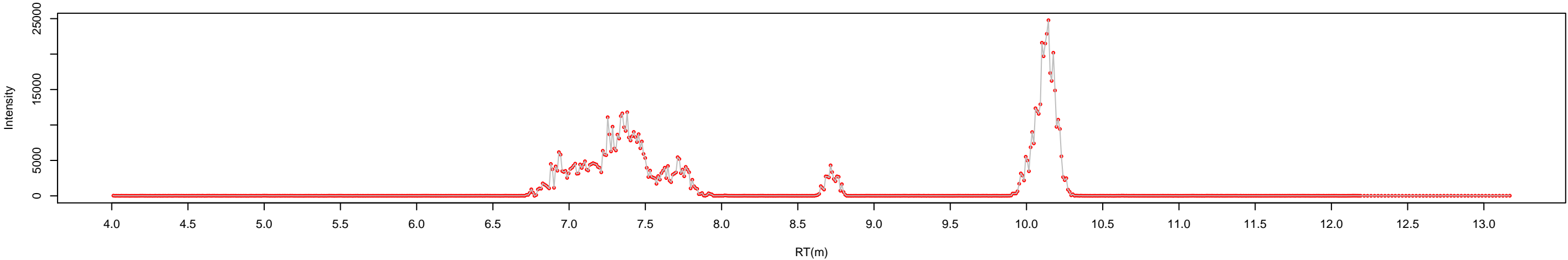


peak area | window size:3   BLine: yes

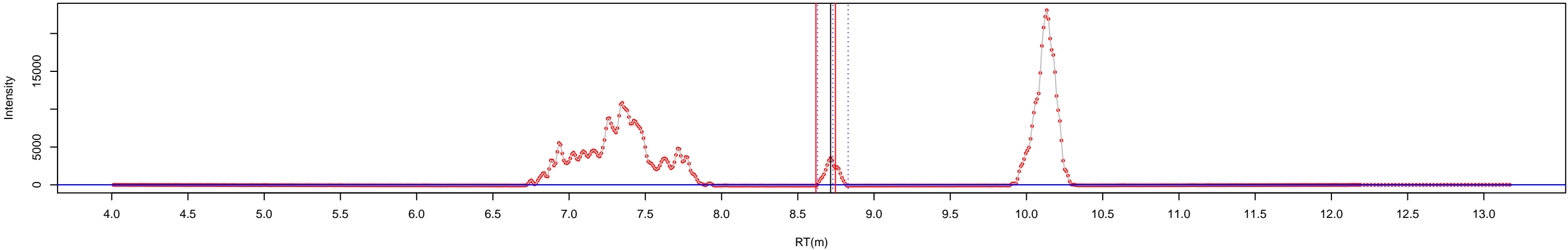




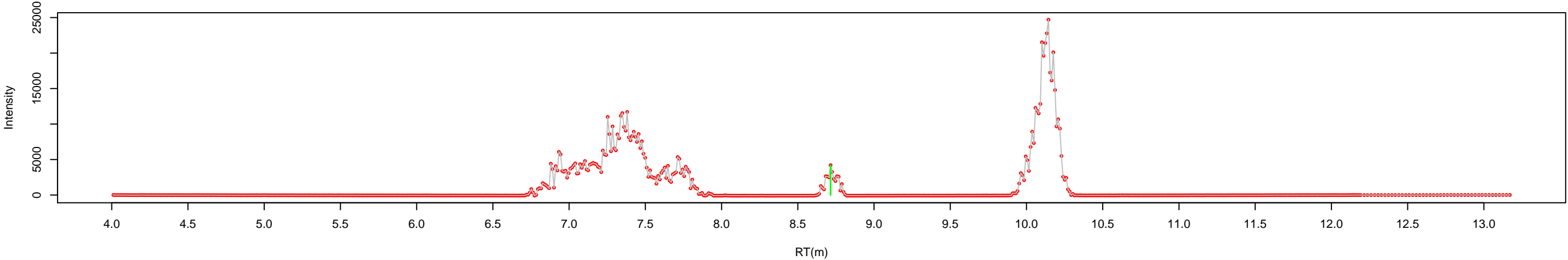
raw chromatogram | batch: 2   sample: std7-10ul-2   conc: 0.2   function: 2   mass: 514.35>79.93



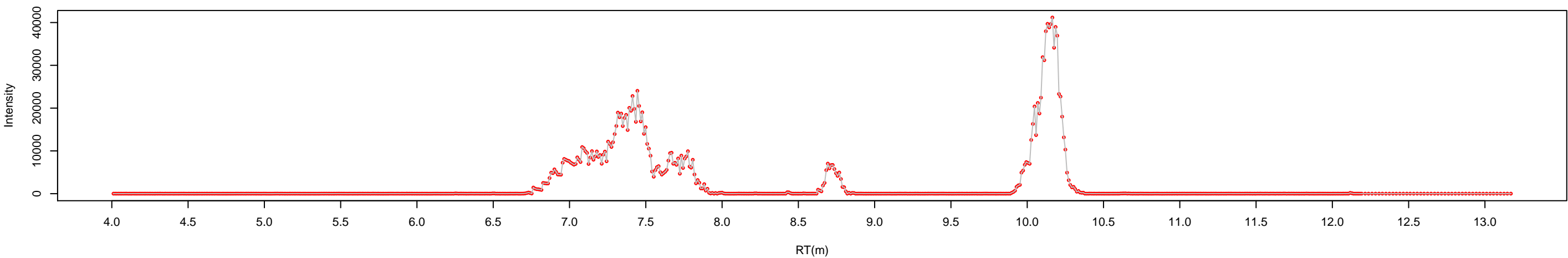
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 2   BLine: yes



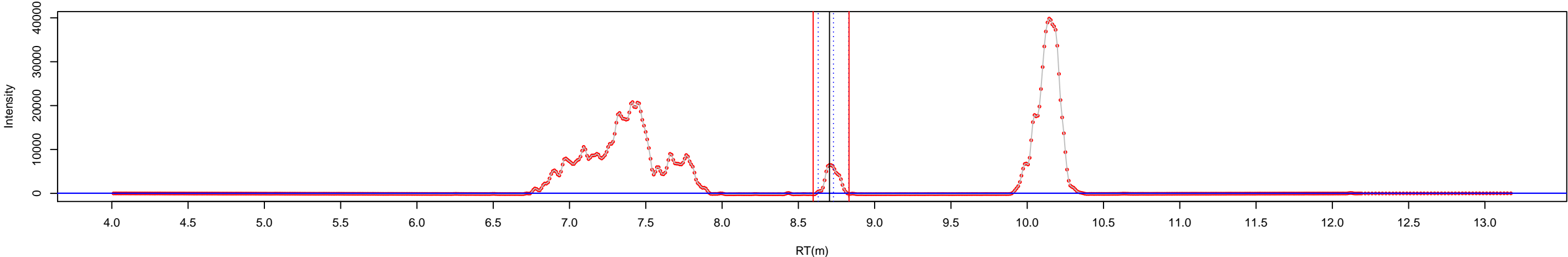
peak area | window size:3   BLine: yes



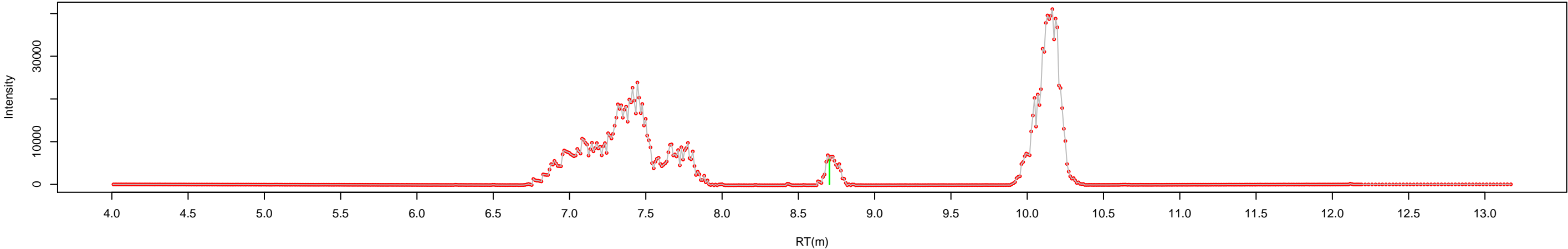
raw chromatogram | batch: 2   sample: std8-10ul-2   conc: 0.5   function: 2   mass: 514.35>79.93



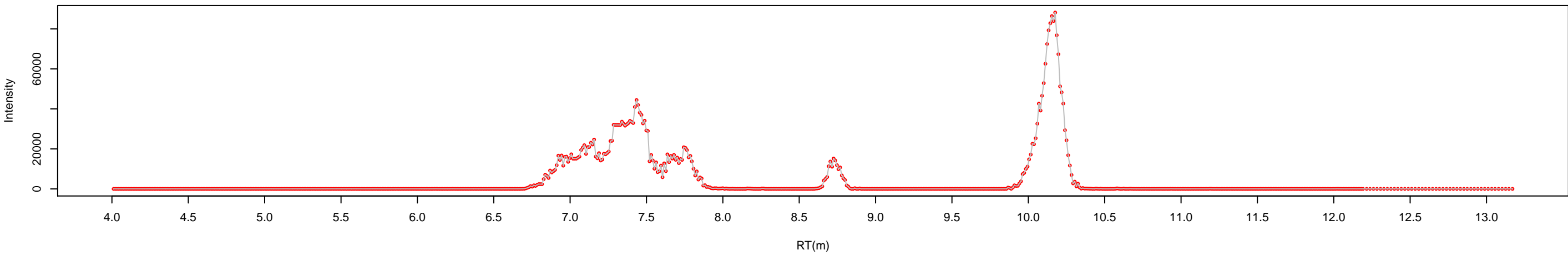
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 4.86   BLine: yes



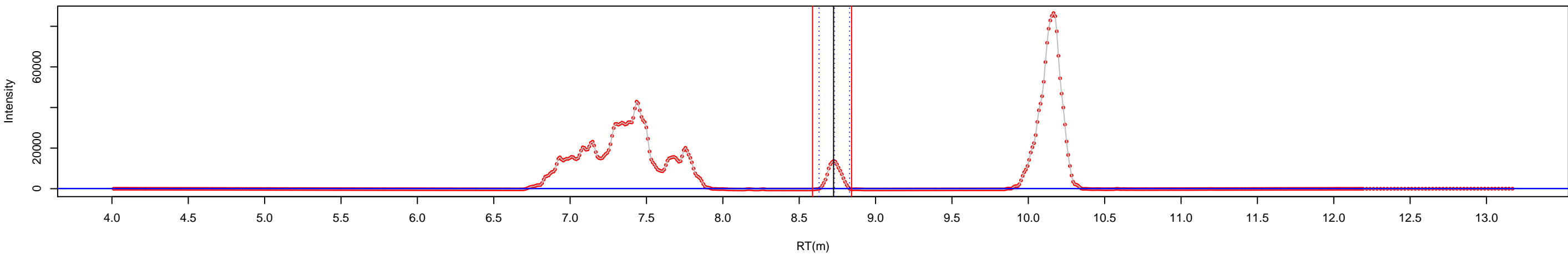
peak area | window size:3   BLine: yes



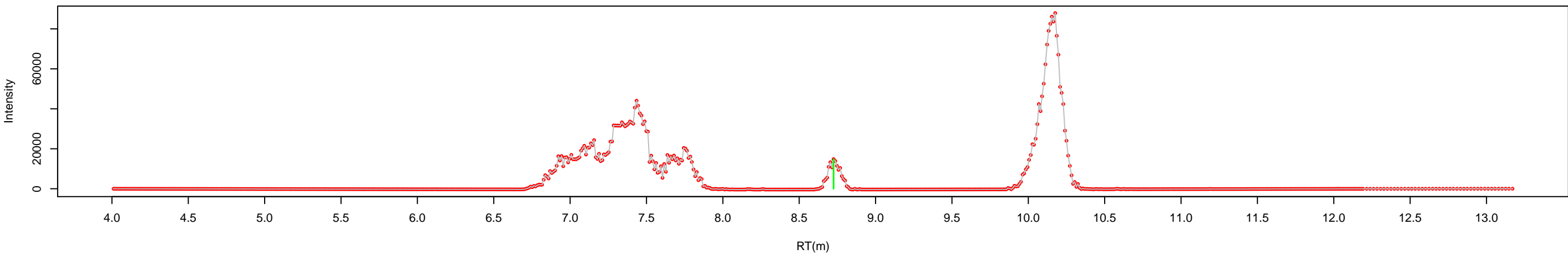
raw chromatogram | batch: 2 sample: std9-10ul-2 conc: 1 function: 2 mass: 514.35>79.93



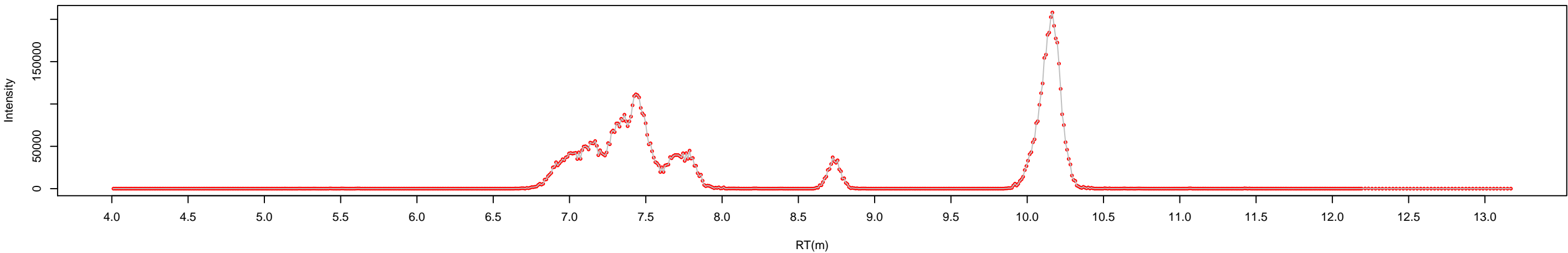
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 12.6 BLine: yes



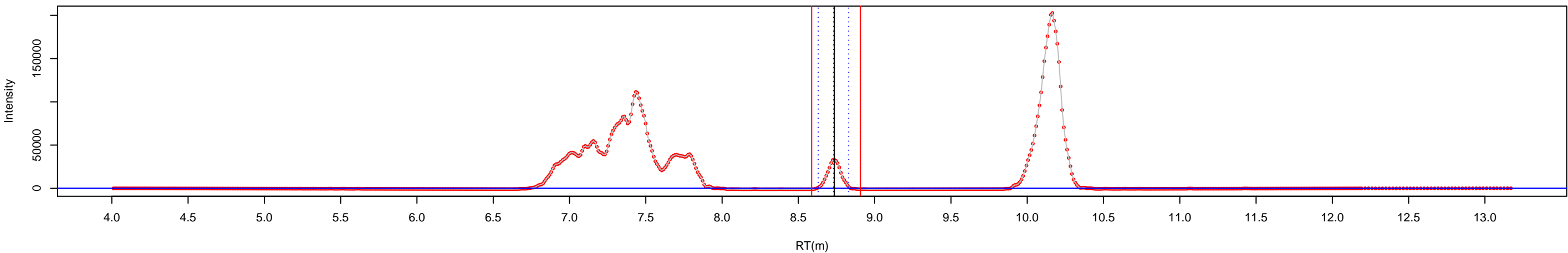
peak area | window size:3 BLine: yes



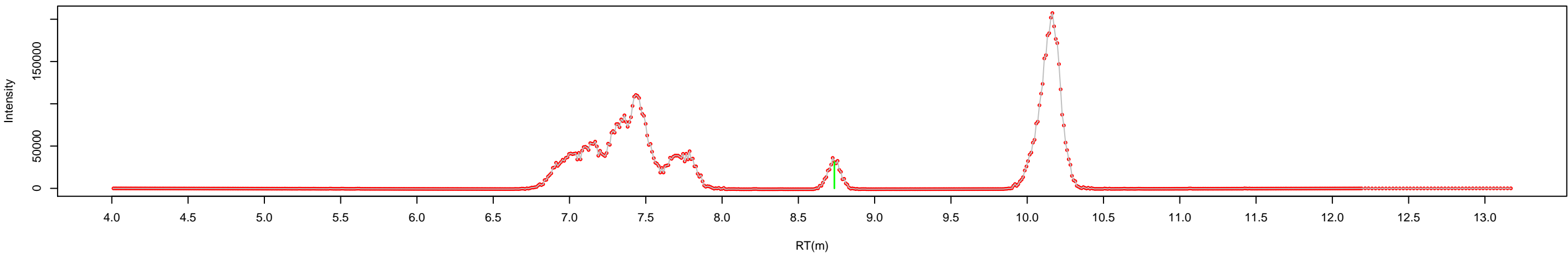
raw chromatogram | batch: 2 sample: std10-10ul-2 conc: 2 function: 2 mass: 514.35>79.93



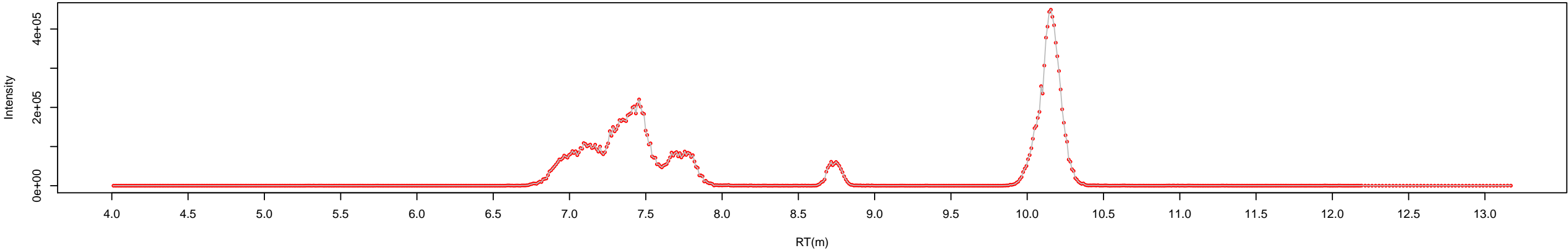
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 10.6 BLine: yes



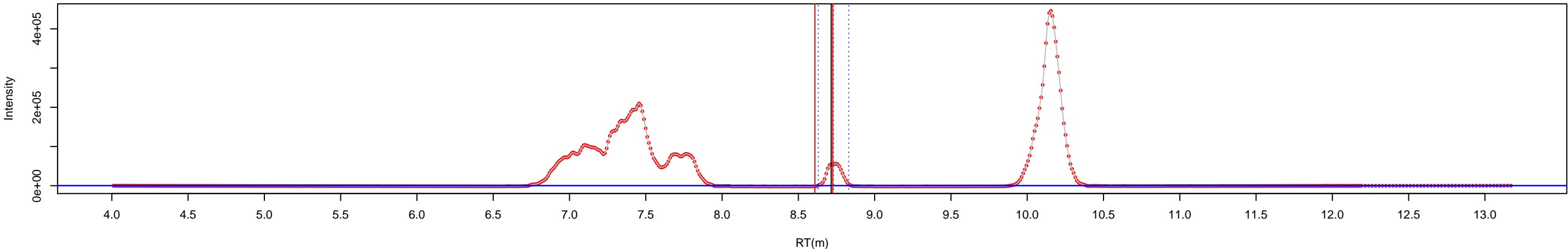
peak area | window size:3 BLine: yes



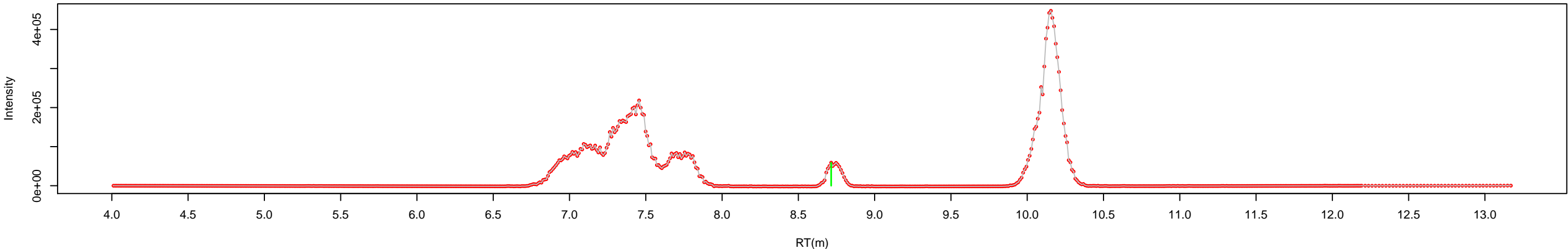
raw chromatogram | batch: 2   sample: std11-10ul-2-1   conc: 5   function: 2   mass: 514.35>79.93



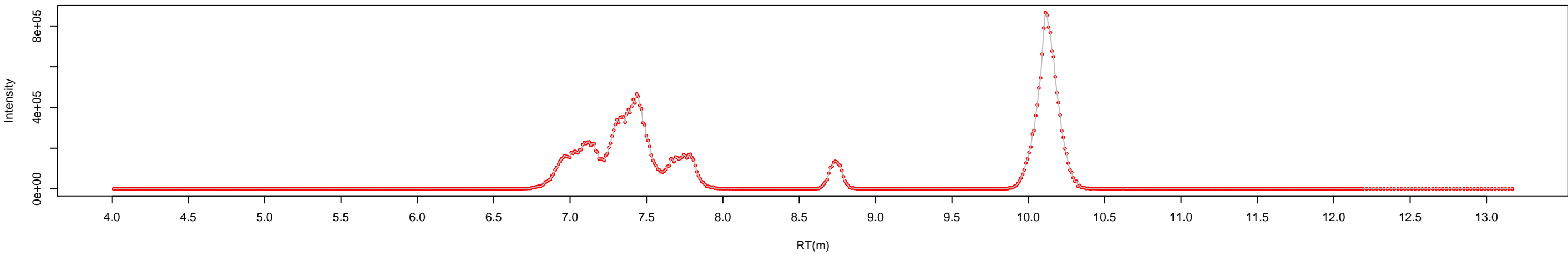
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 39.4   BLine: yes



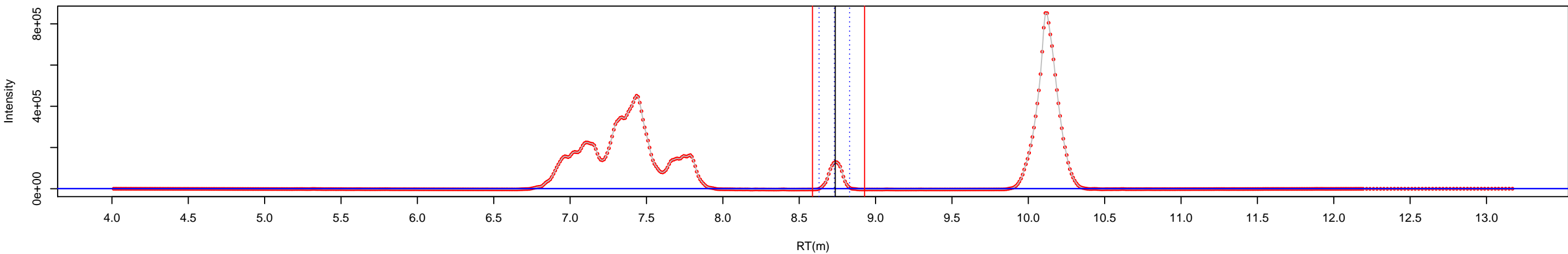
peak area | window size:3   BLine: yes



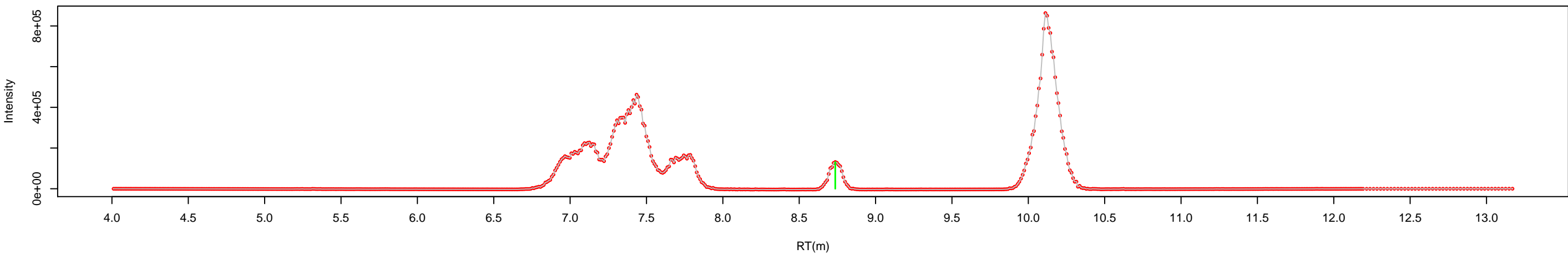
raw chromatogram | batch: 2 sample: std12-10ul-2-1 conc: 10 function: 2 mass: 514.35>79.93



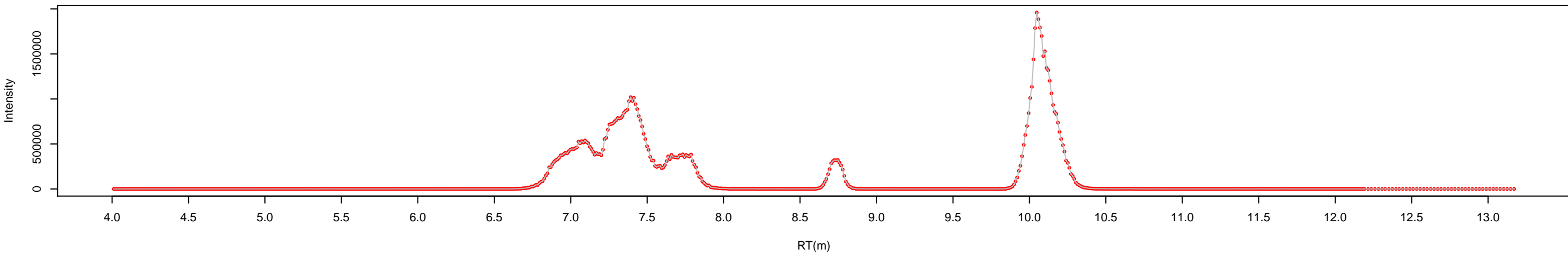
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 105 BLine: yes



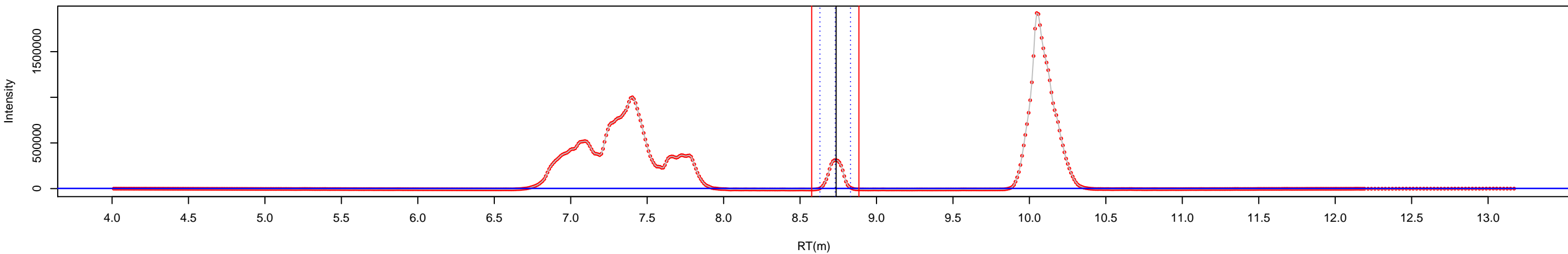
peak area | window size:3 BLine: yes



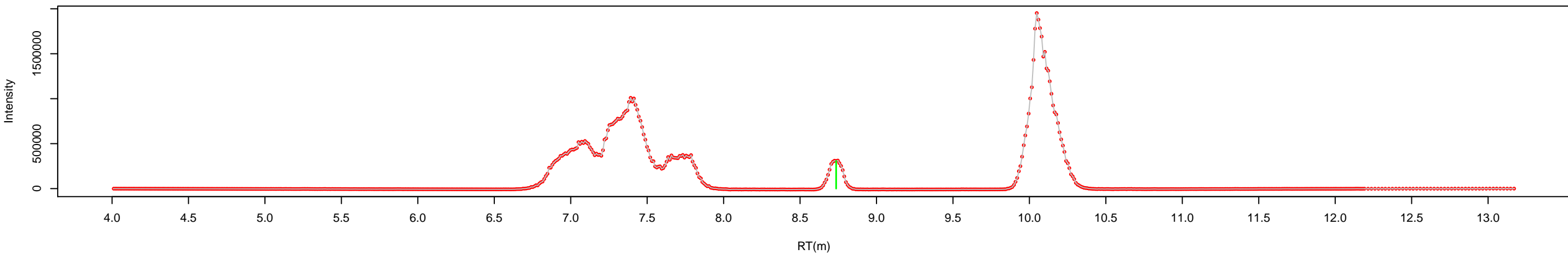
raw chromatogram | batch: 2   sample: std13-10ul-2-1   conc: 20   function: 2   mass: 514.35>79.93



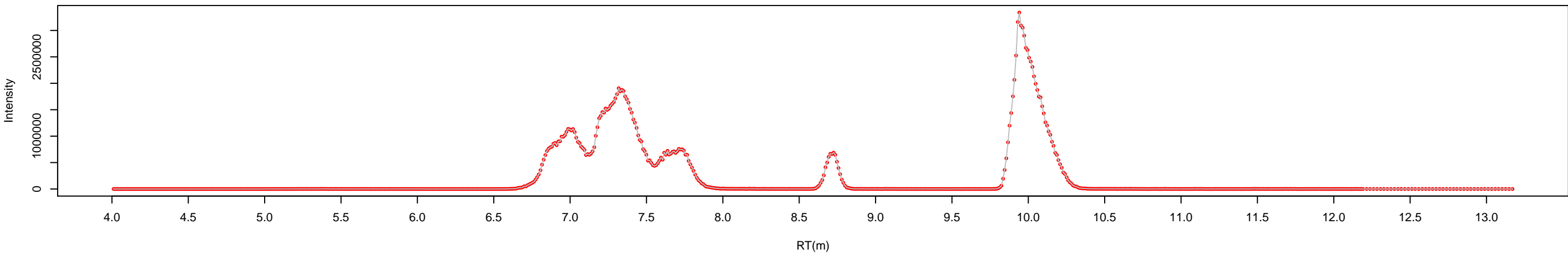
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 384   BLine: yes



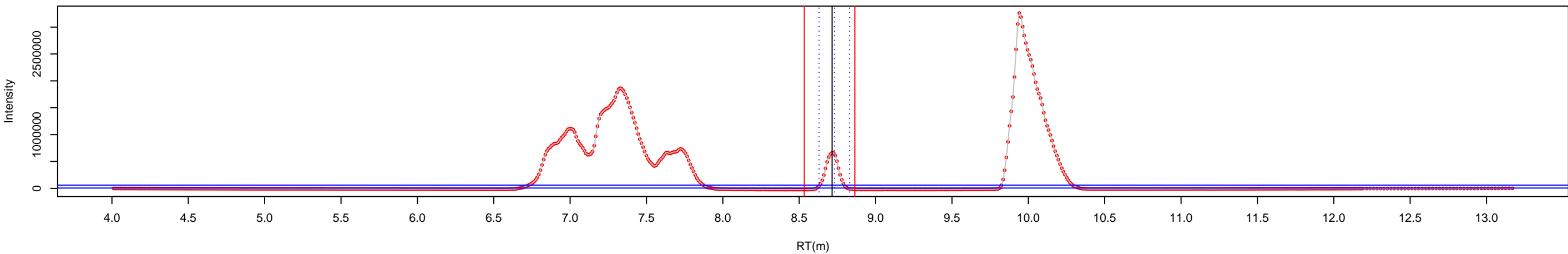
peak area | window size:3   BLine: yes



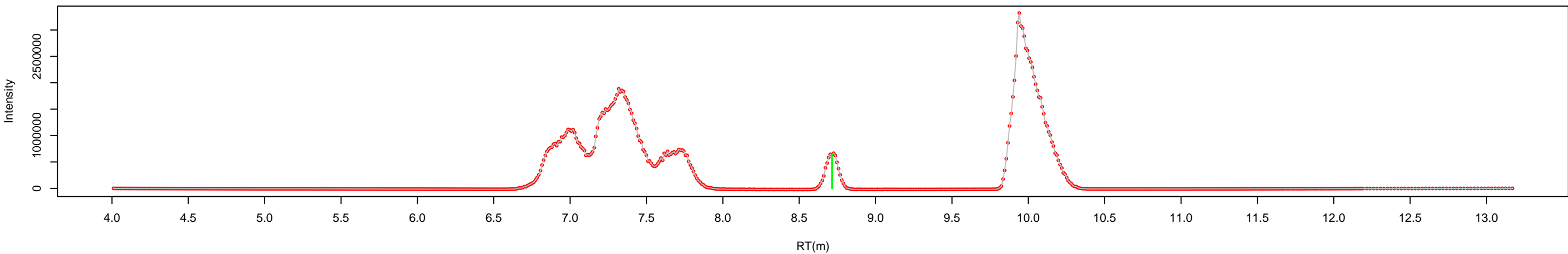
raw chromatogram | batch: 2 sample: std14-10ul-2-1 conc: 50 function: 2 mass: 514.35>79.93



peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 5990 BLine: yes

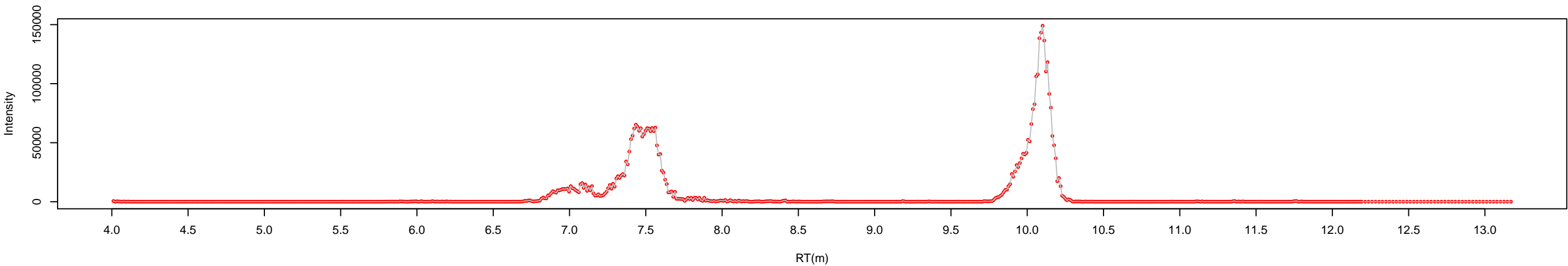


peak area | window size:3 BLine: yes

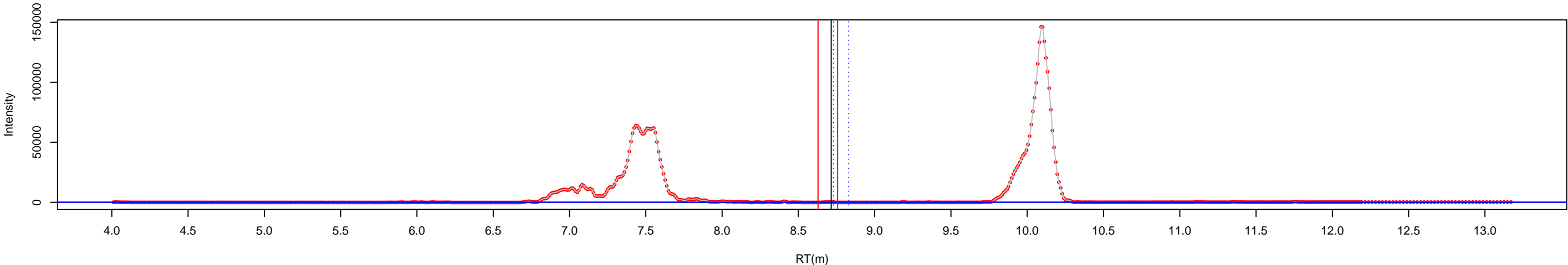




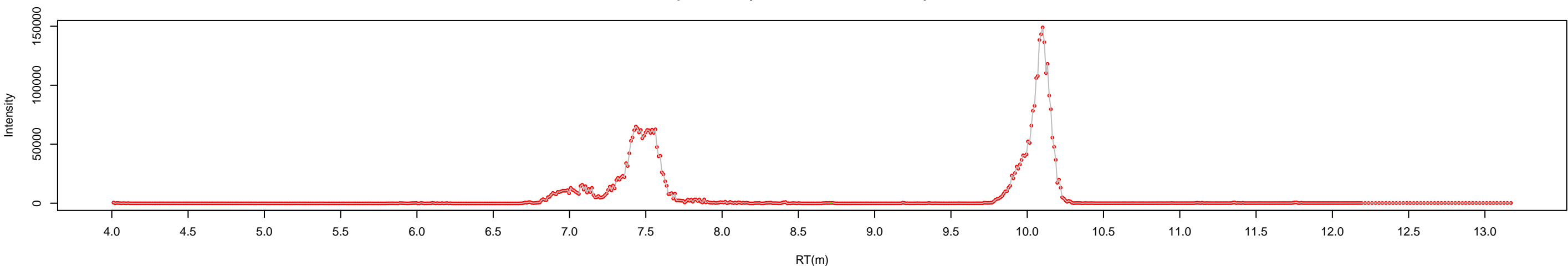
raw chromatogram | batch: 3 sample: cotn65 conc: NA function: 2 mass: 514.35>79.93



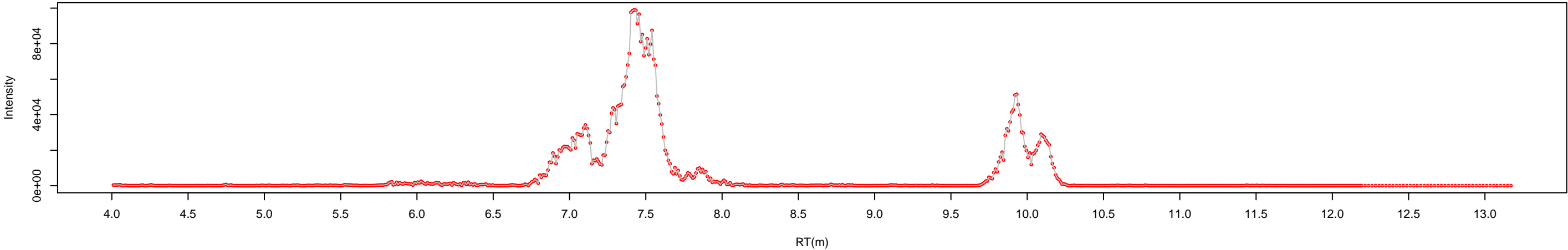
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 13.2 BLine: yes



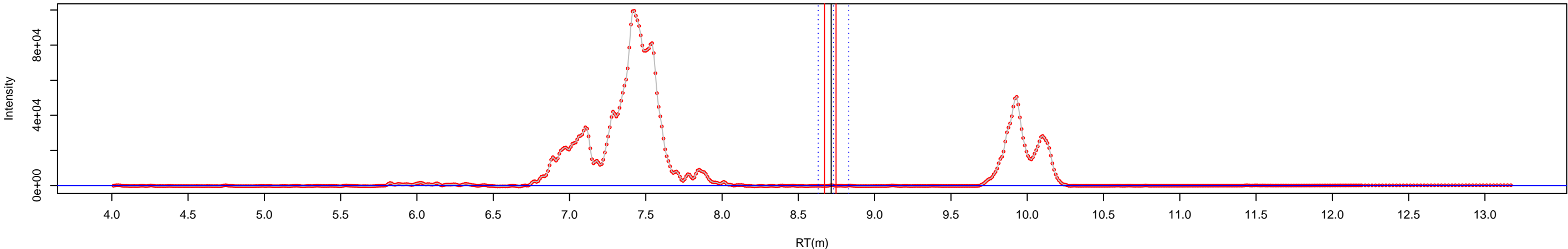
peak area | window size:3 BLine: yes



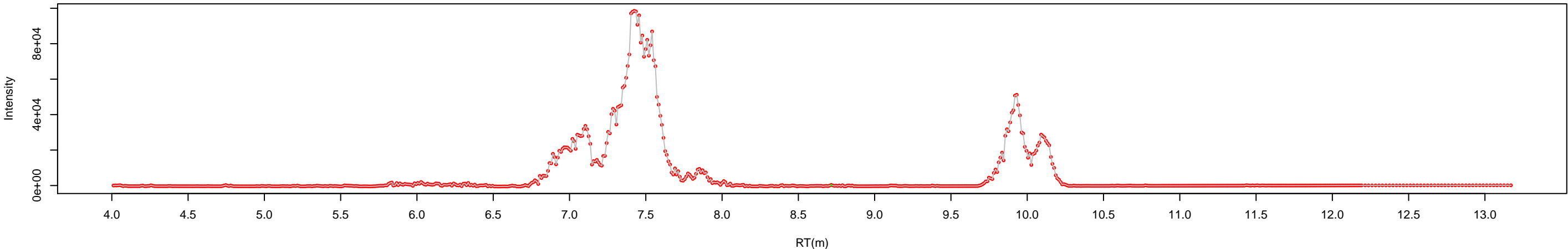
raw chromatogram | batch: 3 sample: cotn66 conc: NA function: 2 mass: 514.35>79.93



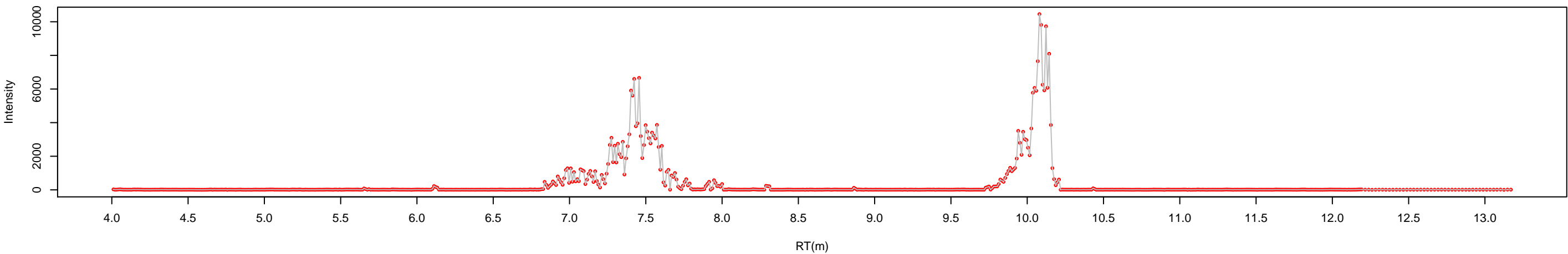
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 3.96 BLine: yes



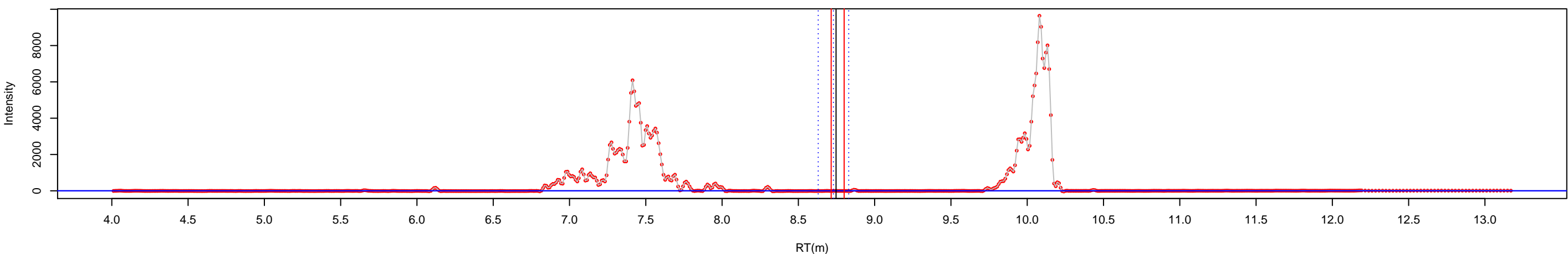
peak area | window size:3 BLine: yes



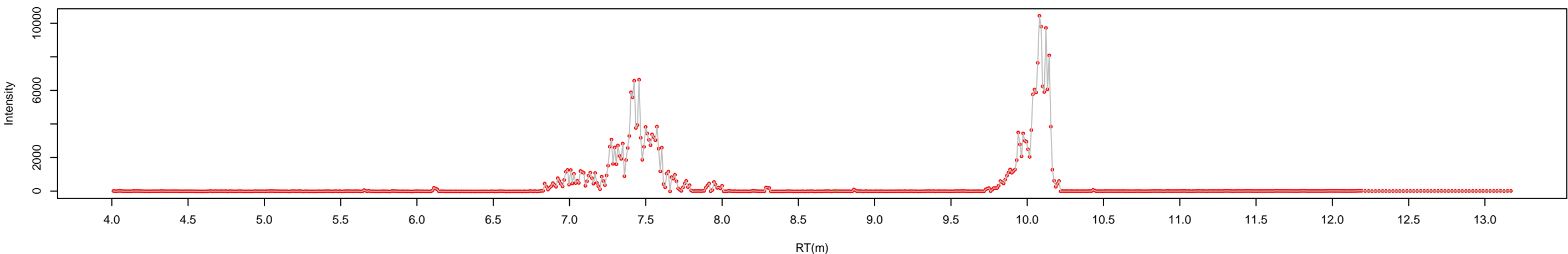
raw chromatogram | batch: 3 sample: cotn67 conc: NA function: 2 mass: 514.35>79.93



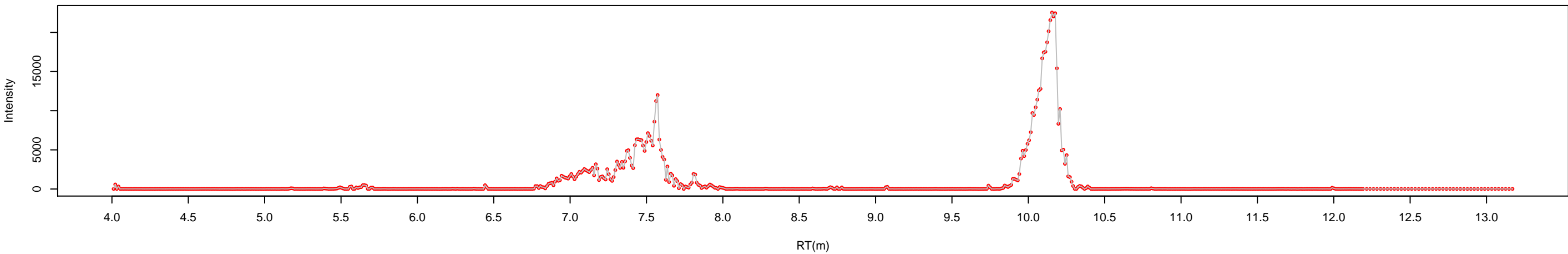
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.34 BLine: yes



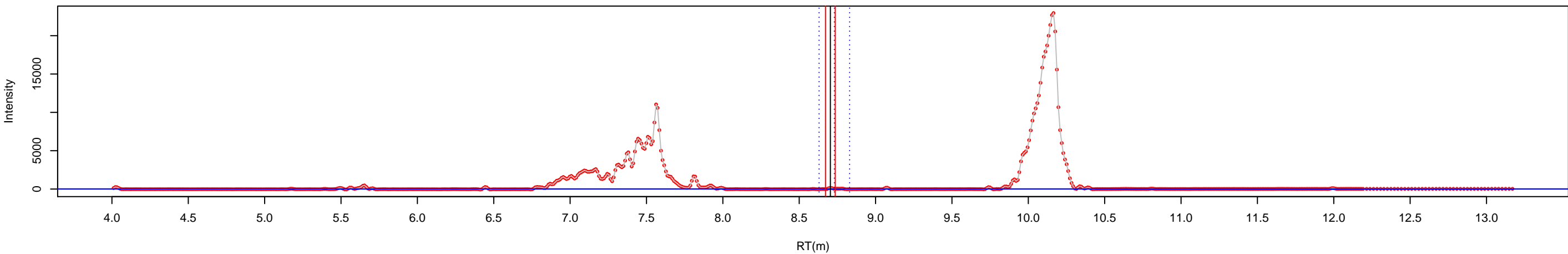
peak area | window size:3 BLine: yes



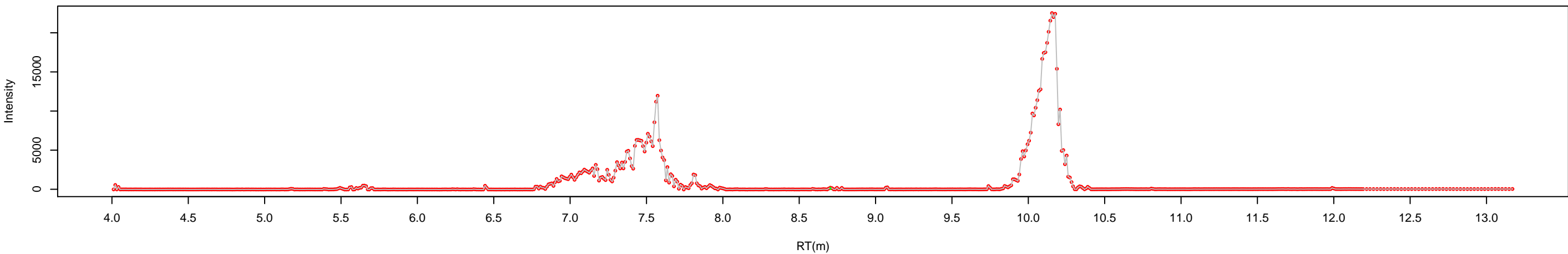
raw chromatogram | batch: 3 sample: cotn195 conc: NA function: 2 mass: 514.35>79.93



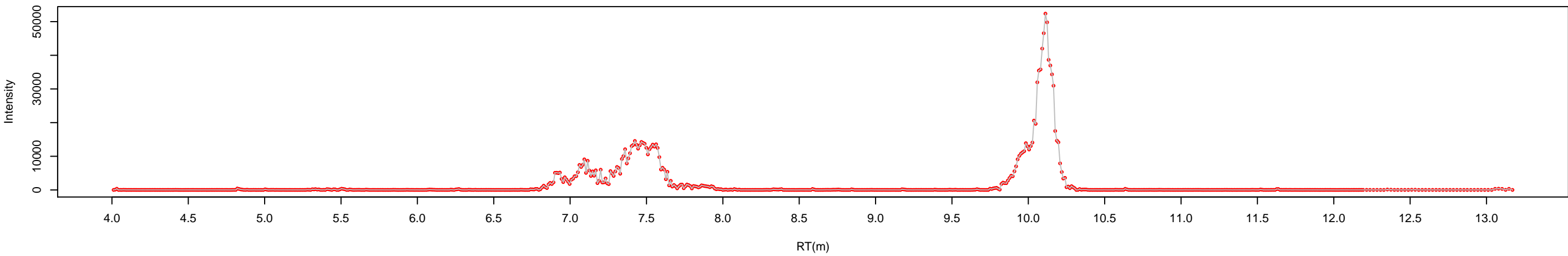
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 2.02 BLine: yes



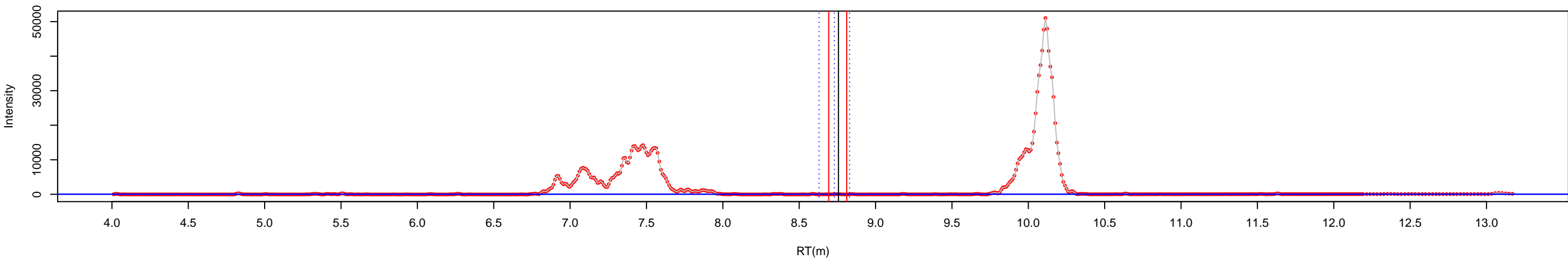
peak area | window size:3 BLine: yes



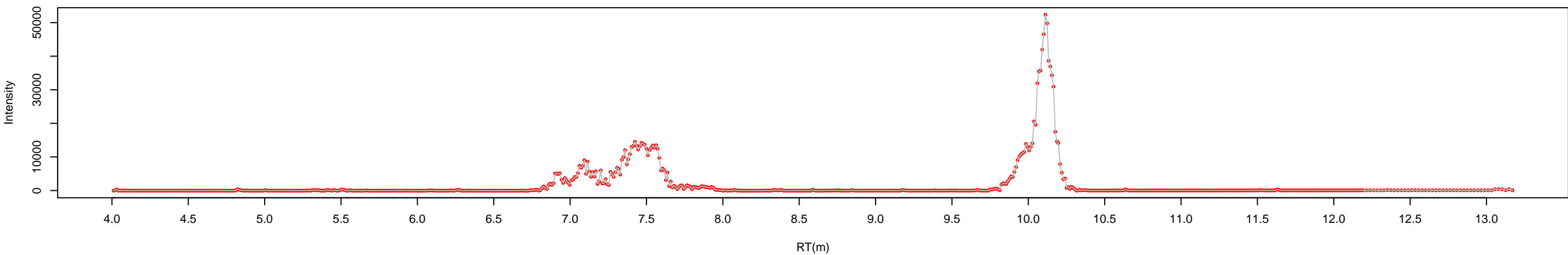
raw chromatogram | batch: 3 sample: cotn196 conc: NA function: 2 mass: 514.35>79.93



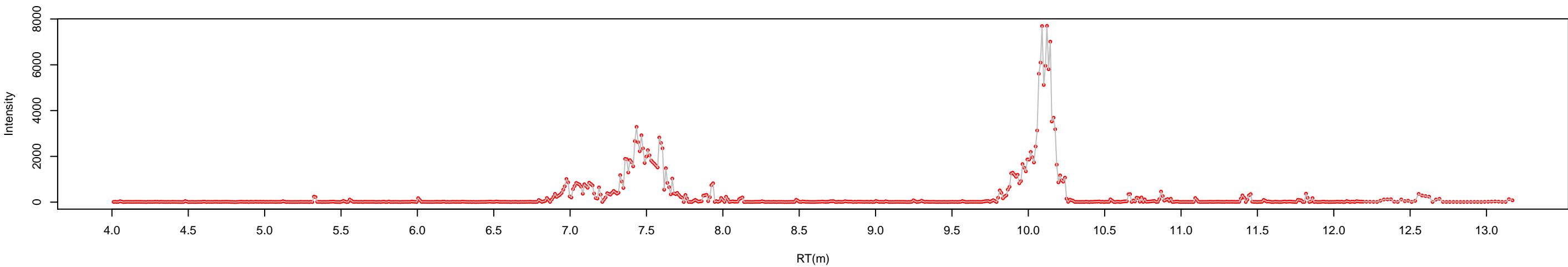
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 3.84 BLine: yes



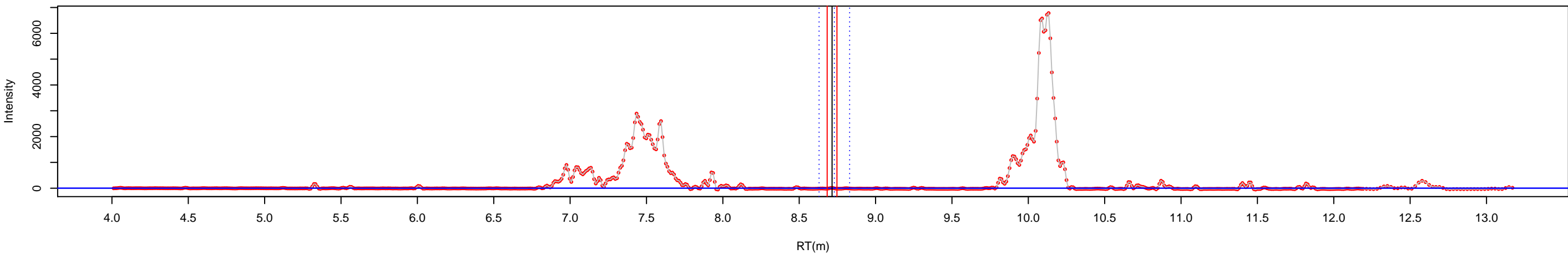
peak area | window size:3 BLine: yes



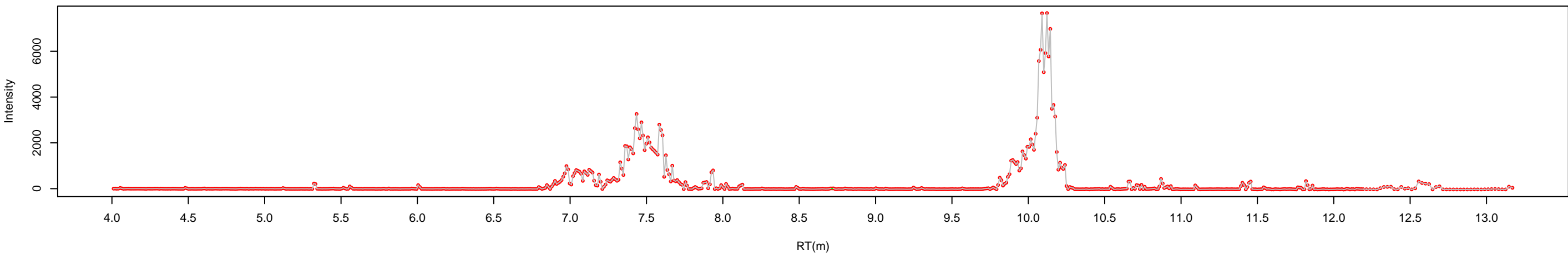
raw chromatogram | batch: 3 sample: cotn197 conc: NA function: 2 mass: 514.35>79.93



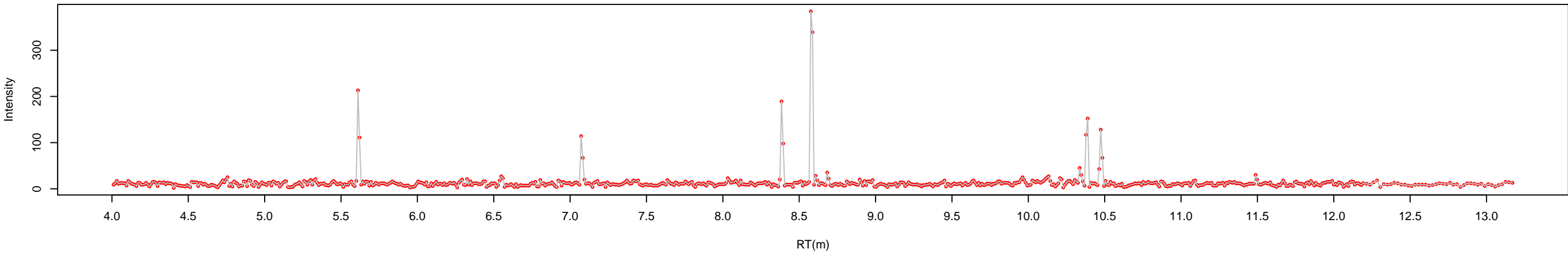
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.348 BLine: yes



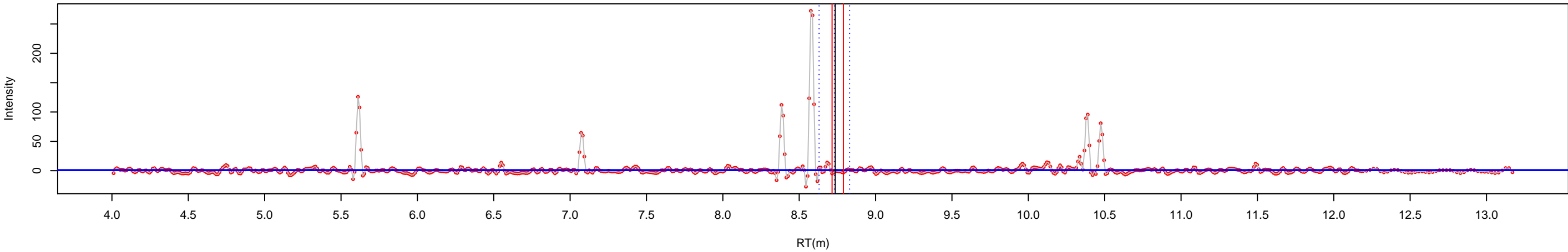
peak area | window size:3 BLine: yes



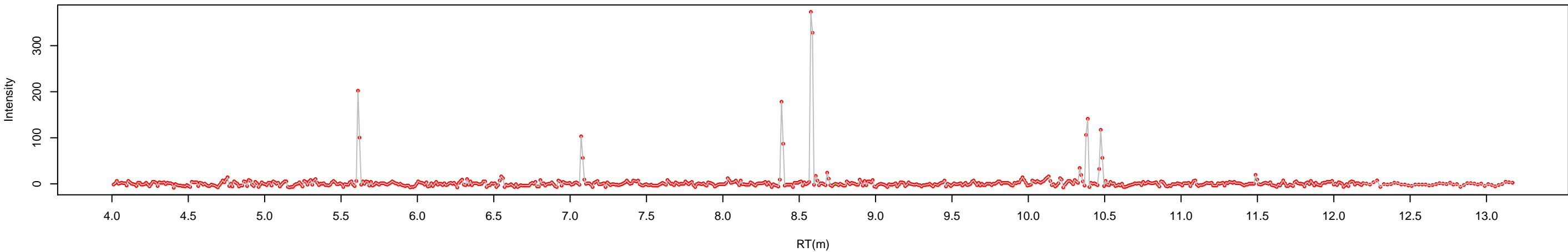
raw chromatogram | batch: 3 sample: std0-10ul-3 conc: 0.001 function: 2 mass: 514.35>79.93



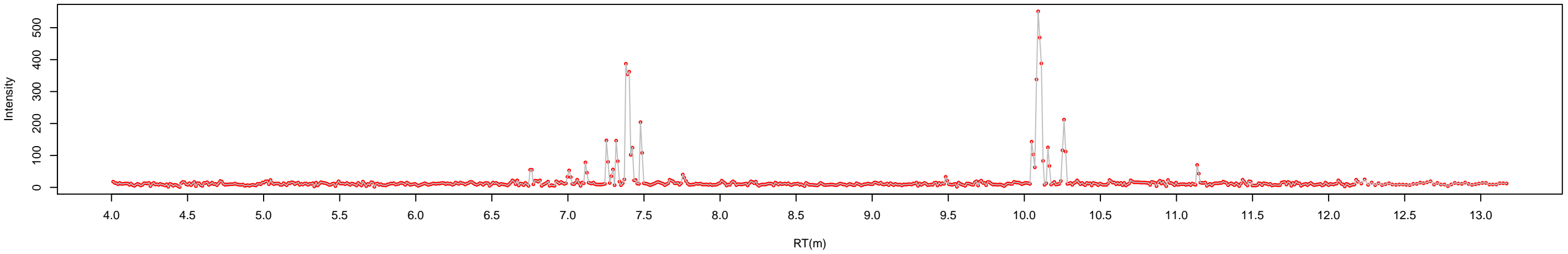
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.174 BLine: yes



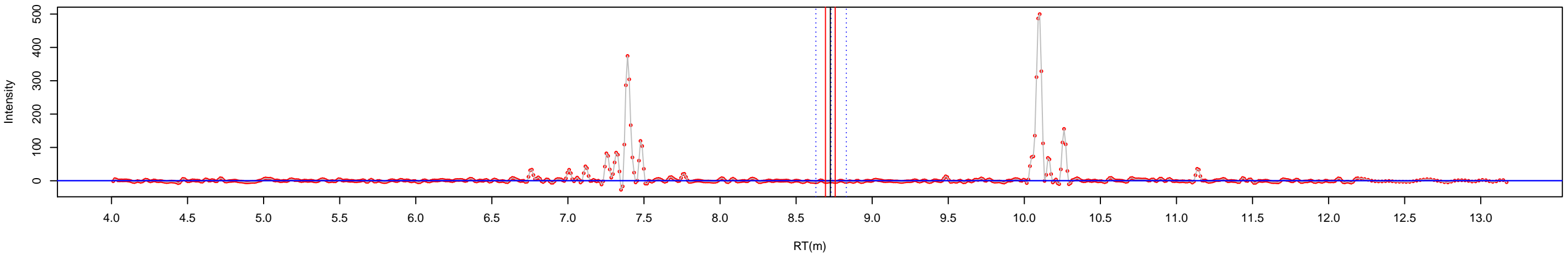
peak area | window size:3 BLine: yes



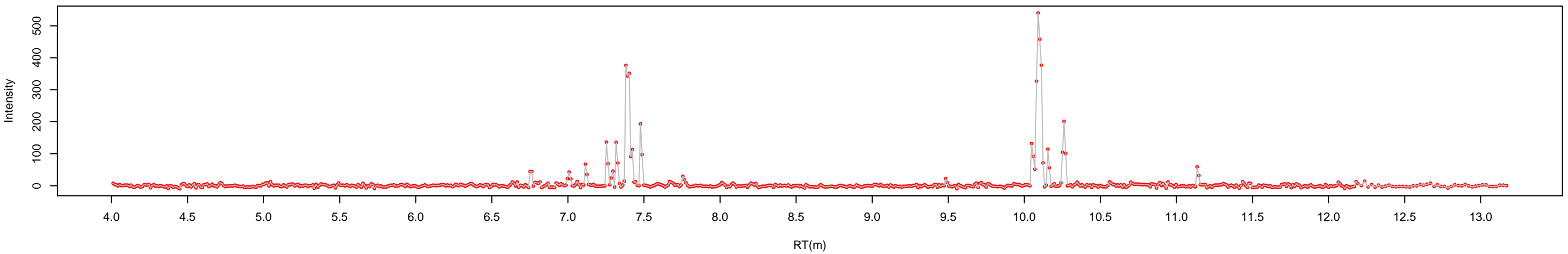
raw chromatogram | batch: 3 sample: std1-10ul-3 conc: 0.002 function: 2 mass: 514.35>79.93



peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.0477 BLine: yes

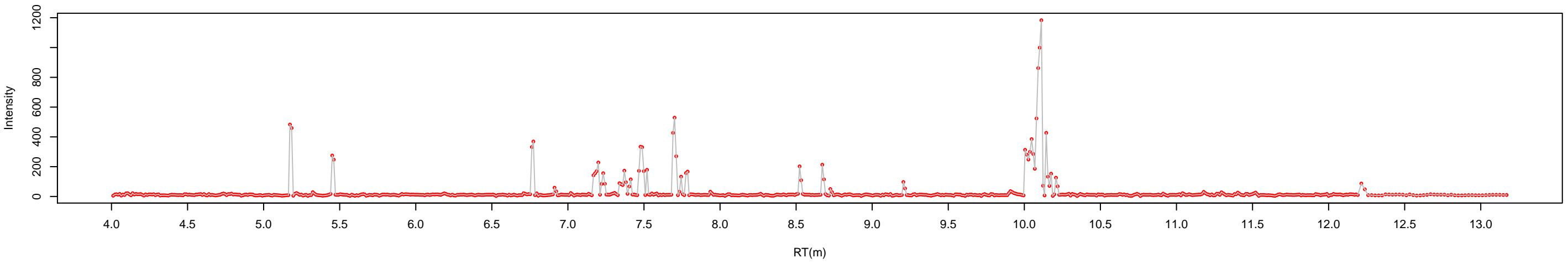


peak area | window size:3 BLine: yes

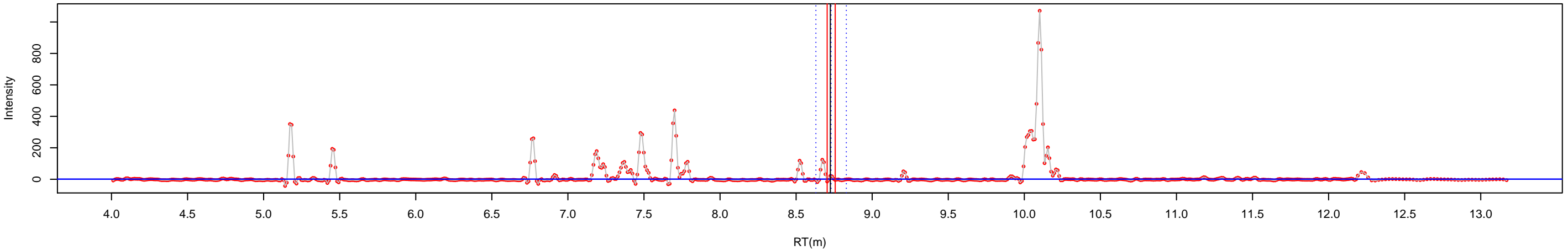




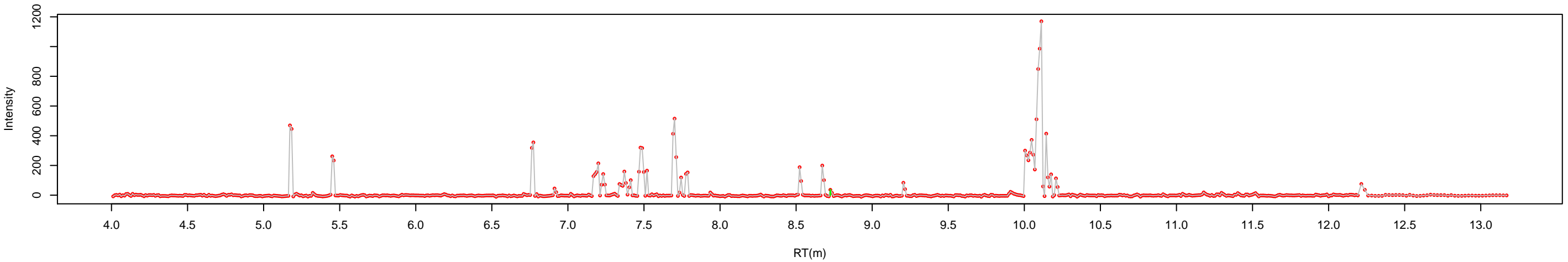
raw chromatogram | batch: 3   sample: std2-10ul-3   conc: 0.005   function: 2   mass: 514.35>79.93



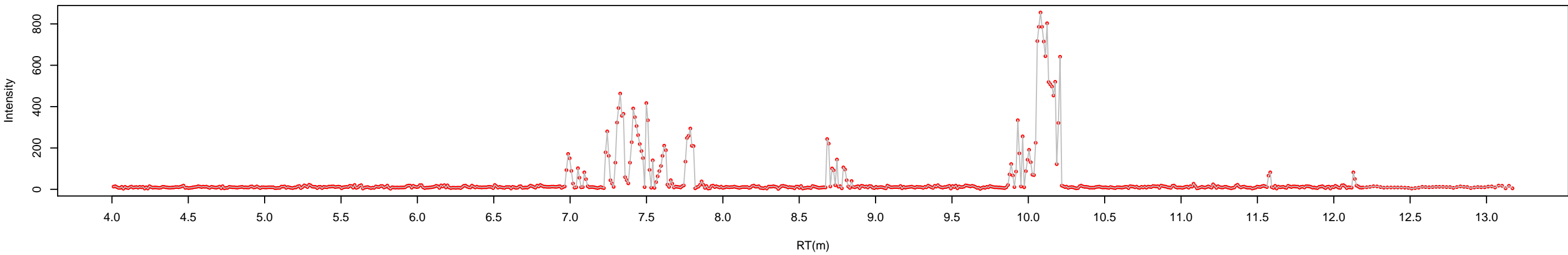
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 0.148   BLine: yes



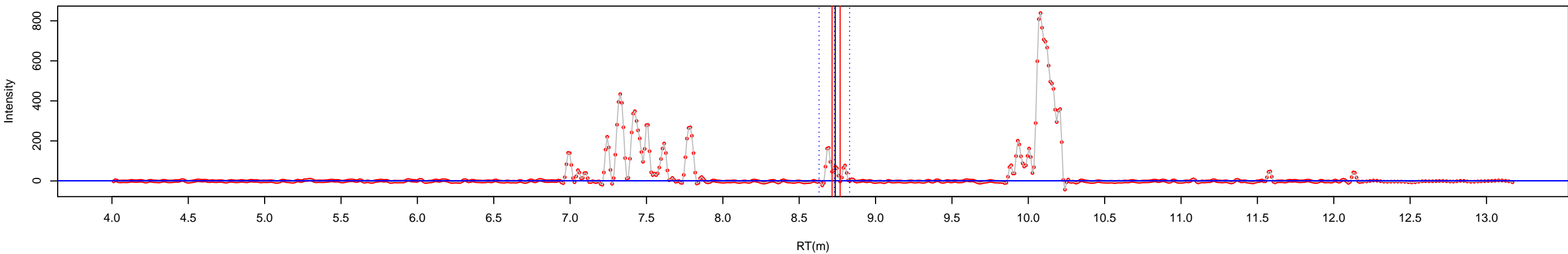
peak area | window size:3   BLine: yes



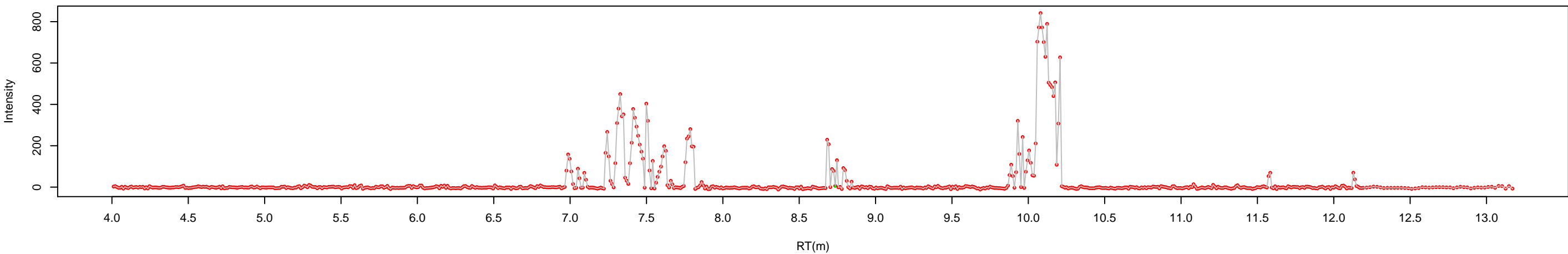
raw chromatogram | batch: 3 sample: std3-10ul-3 conc: 0.01 function: 2 mass: 514.35>79.93



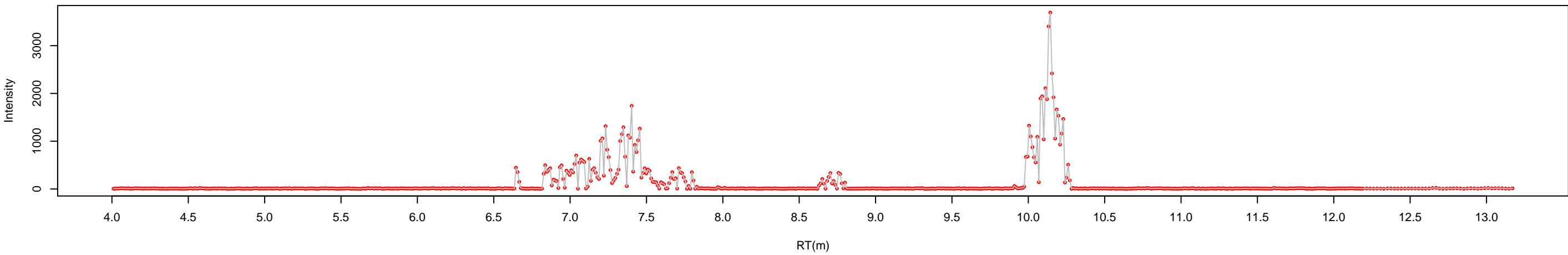
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.144 BLine: yes



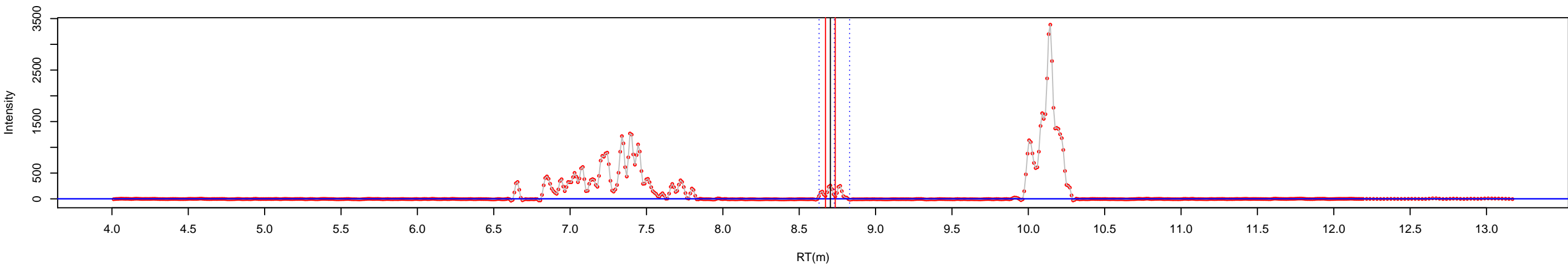
peak area | window size:3 BLine: yes



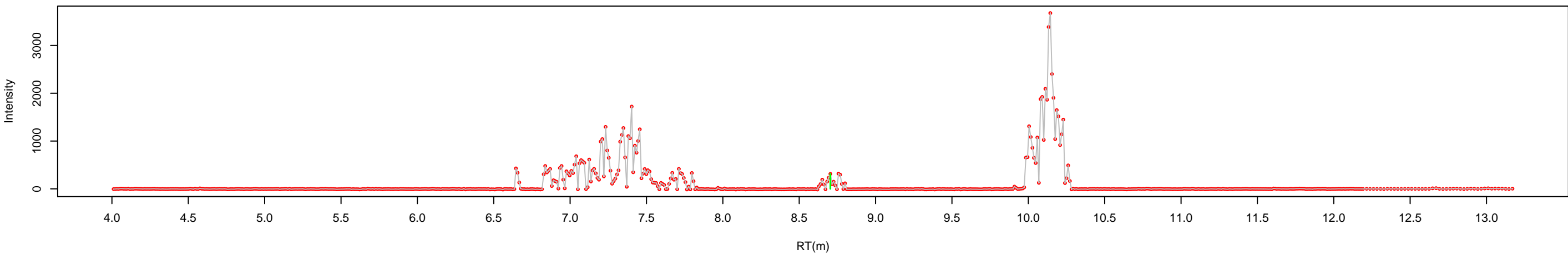
raw chromatogram | batch: 3 sample: std4-10ul-3 conc: 0.02 function: 2 mass: 514.35>79.93



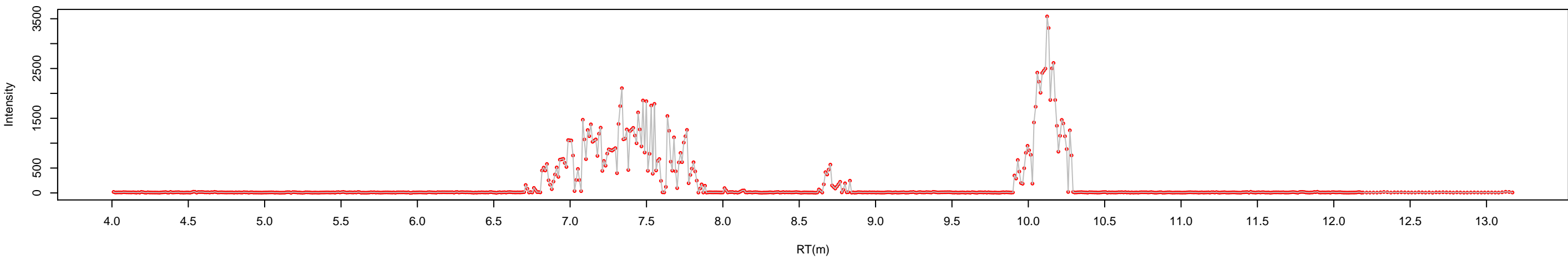
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.15 BLine: yes



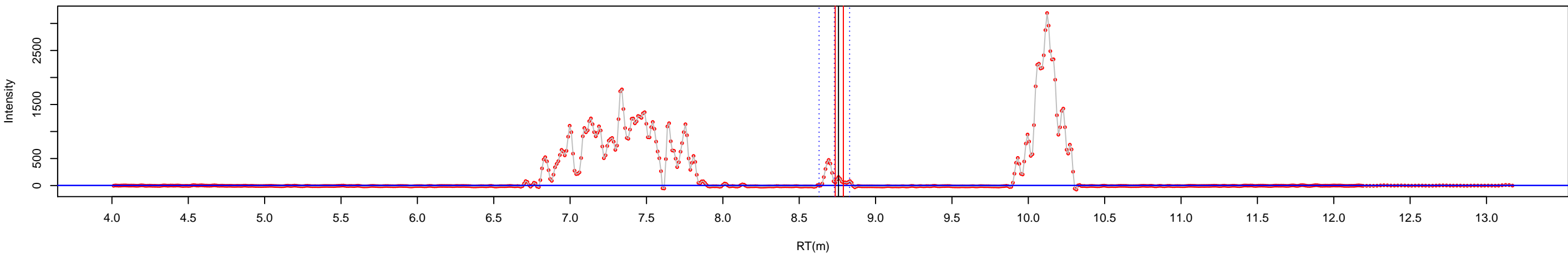
peak area | window size:3 BLine: yes



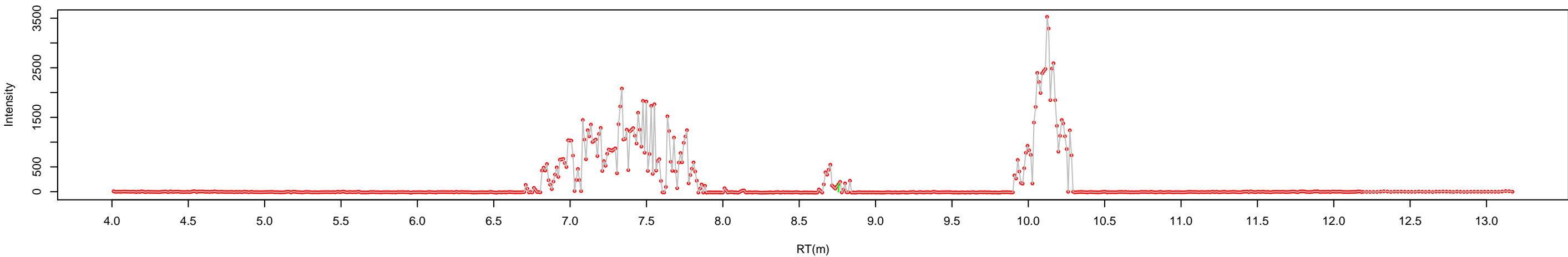
raw chromatogram | batch: 3 sample: std5-10ul-3 conc: 0.05 function: 2 mass: 514.35>79.93



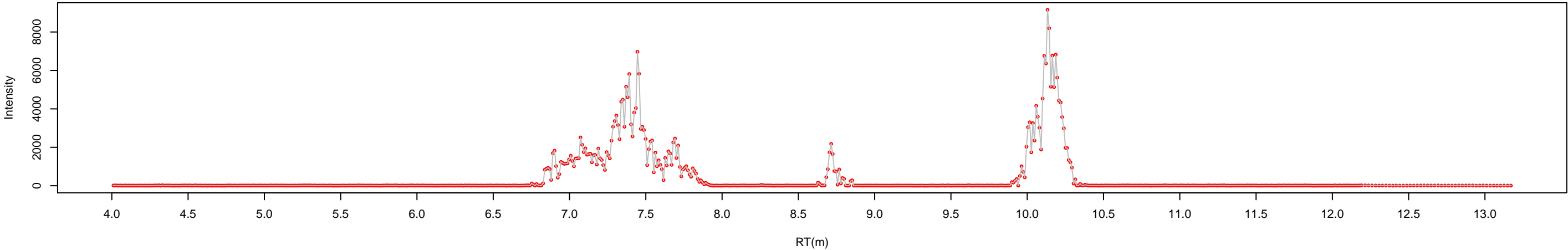
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.397 BLine: yes



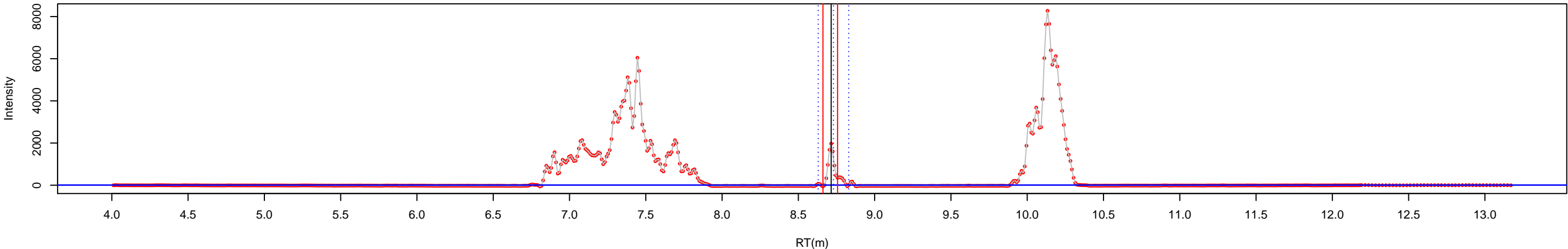
peak area | window size:3 BLine: yes



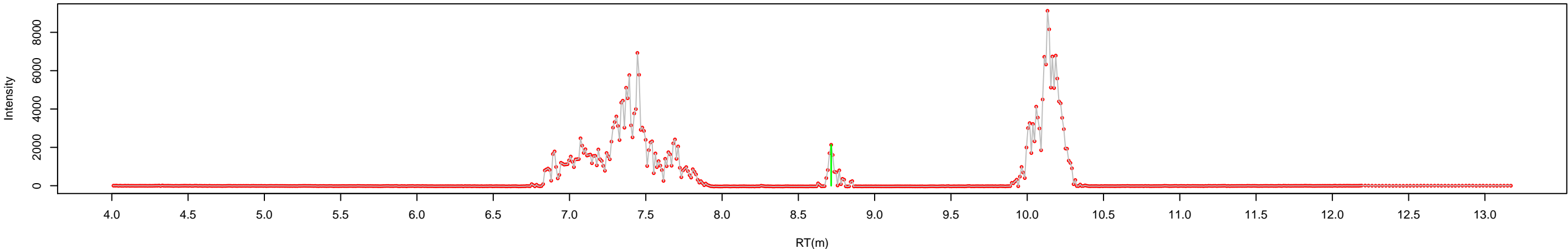
raw chromatogram | batch: 3 sample: std6-10ul-3 conc: 0.1 function: 2 mass: 514.35>79.93



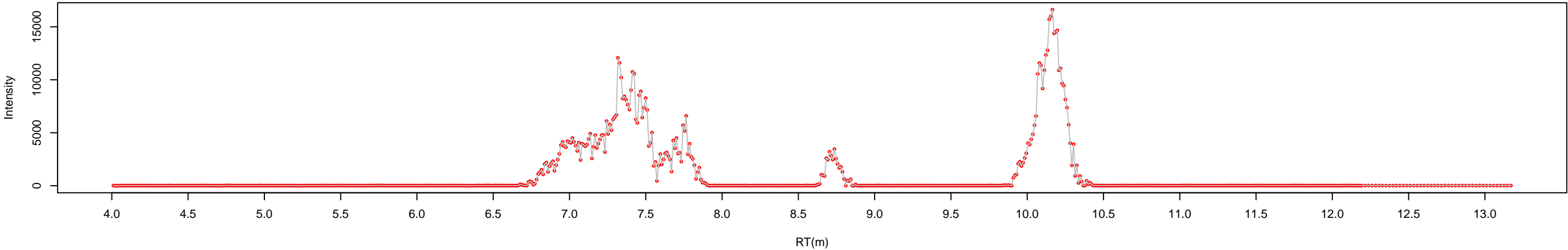
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 1.16 BLine: yes



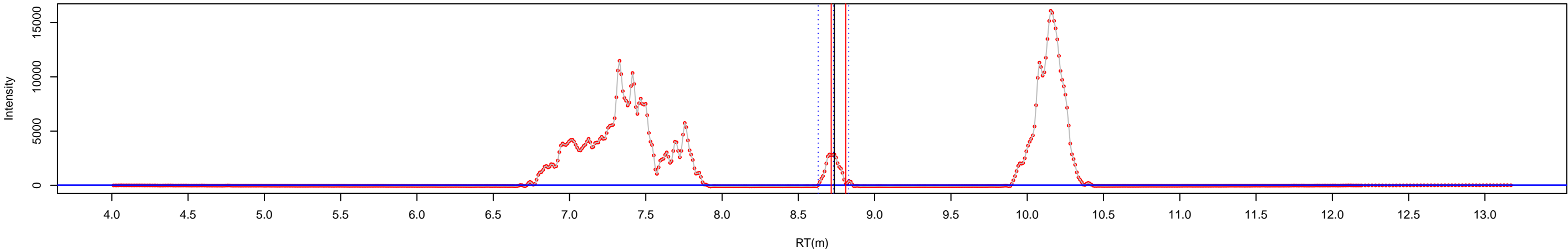
peak area | window size:3 BLine: yes



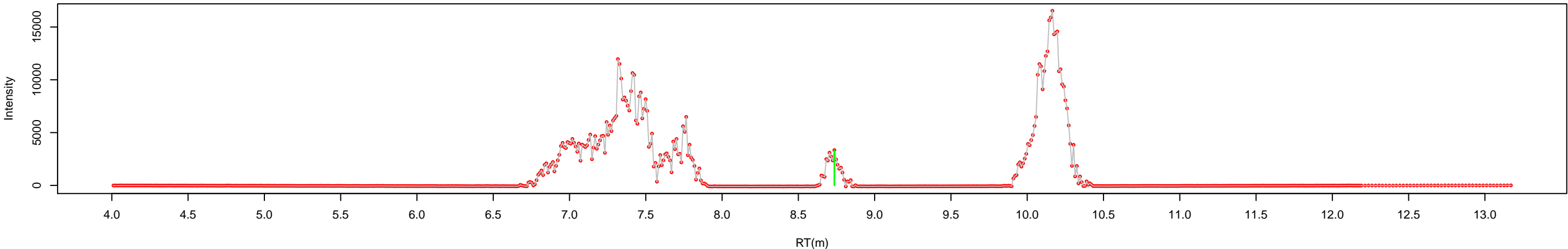
raw chromatogram | batch: 3   sample: std7-10ul-3   conc: 0.2   function: 2   mass: 514.35>79.93



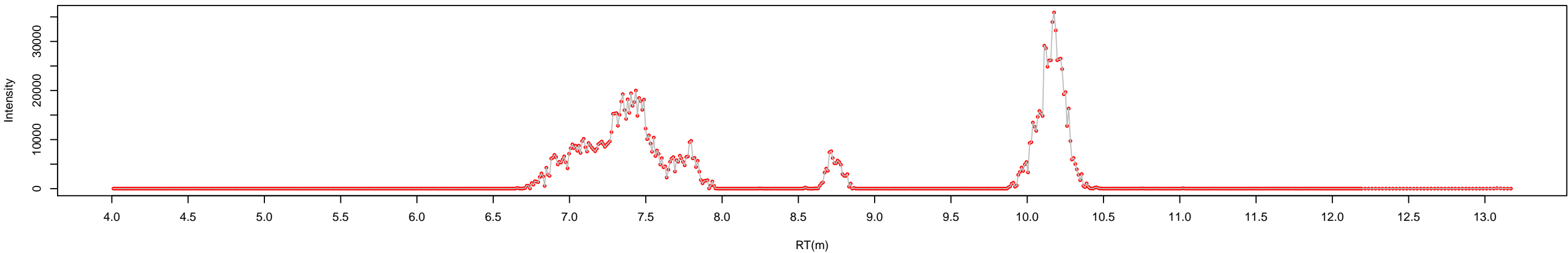
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 2.56   BLine: yes



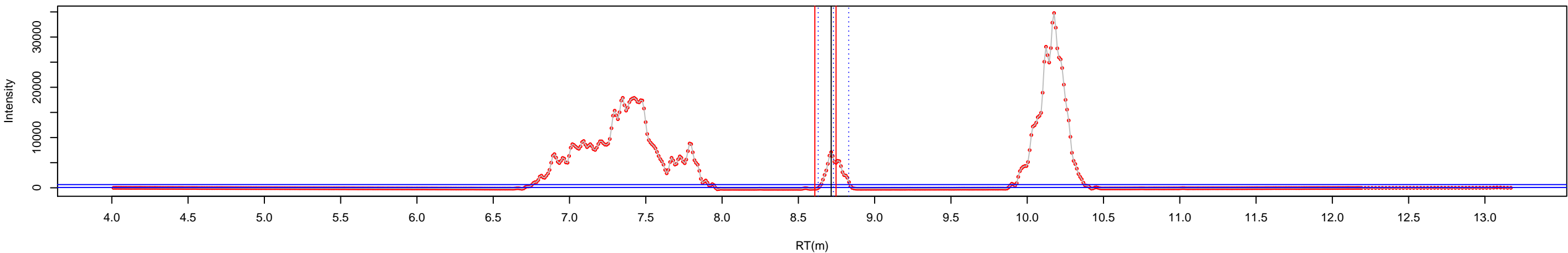
peak area | window size:3   BLine: yes



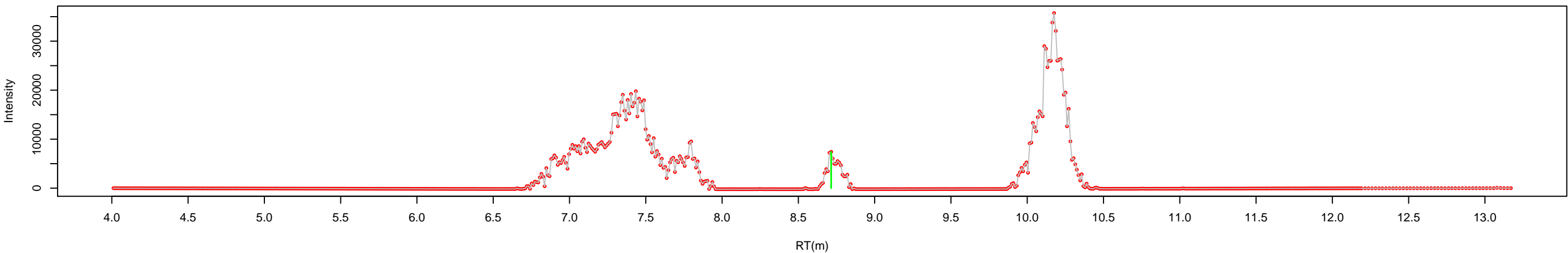
raw chromatogram | batch: 3 sample: std8-10ul-3 conc: 0.5 function: 2 mass: 514.35>79.93



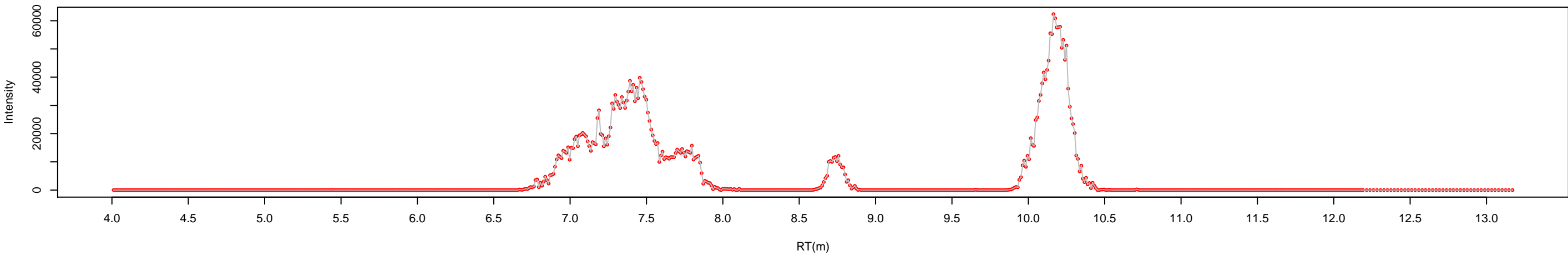
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 64 BLine: yes



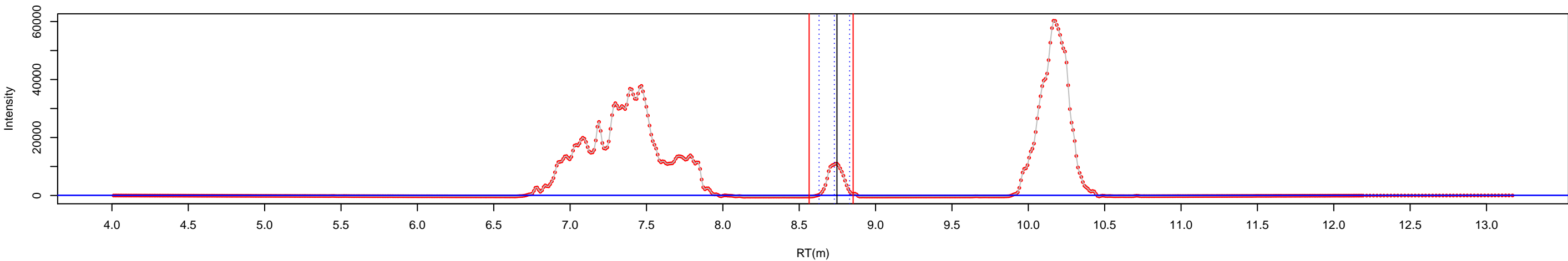
peak area | window size:3 BLine: yes



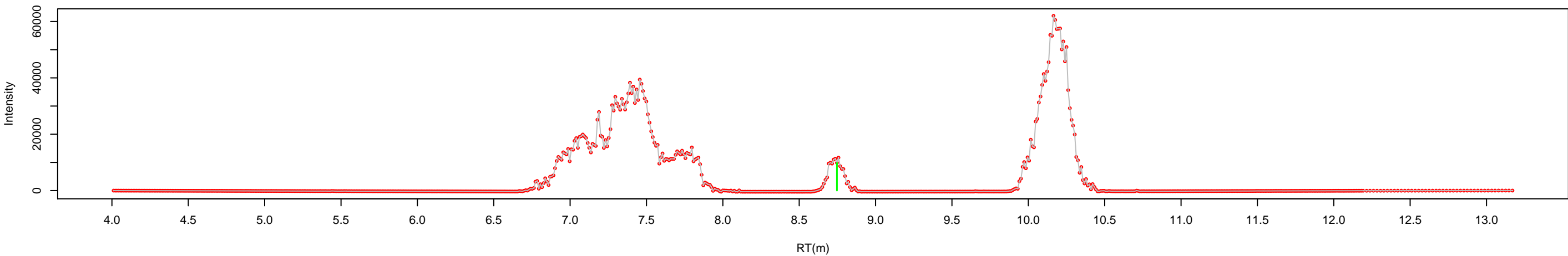
raw chromatogram | batch: 3 sample: std9-10ul-3 conc: 1 function: 2 mass: 514.35>79.93



peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 9.07 BLine: yes

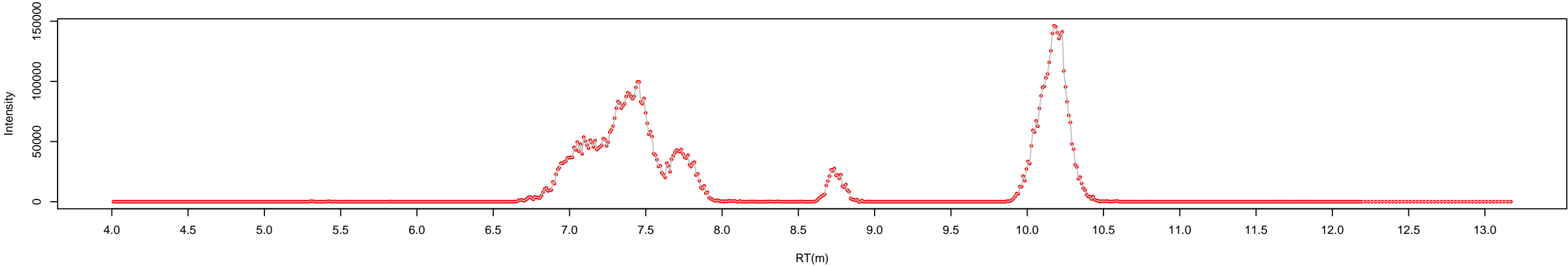


peak area | window size:3 BLine: yes

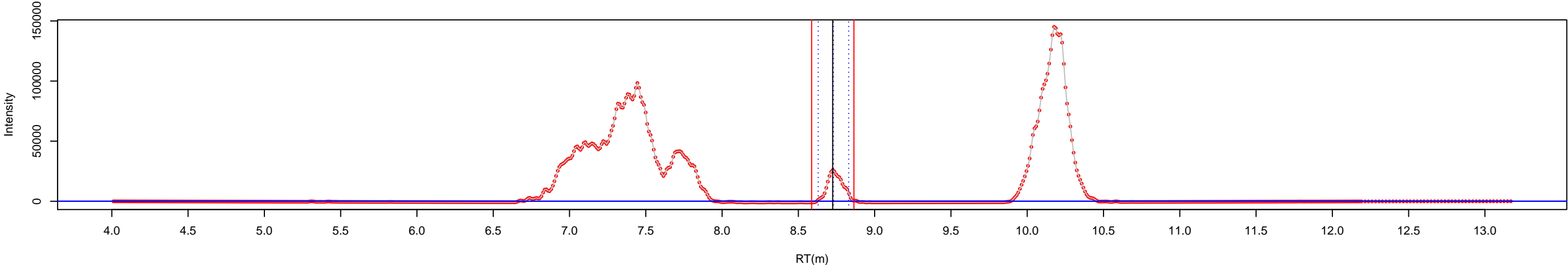




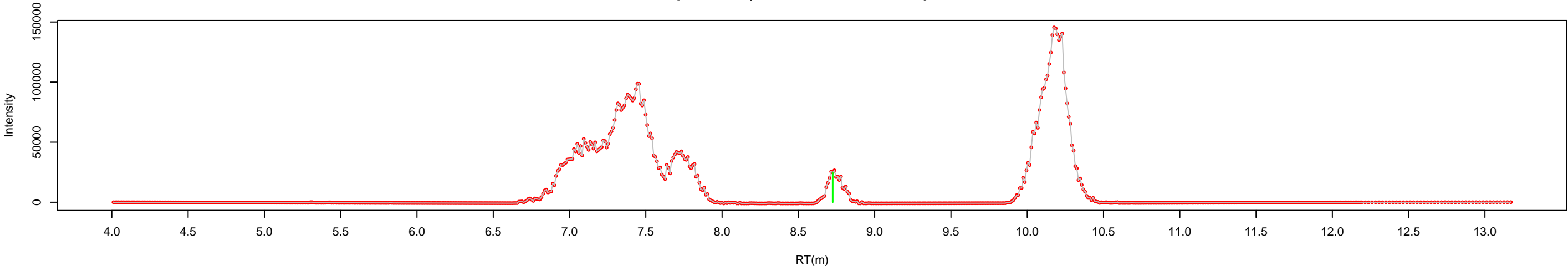
raw chromatogram | batch: 3 sample: std10-10ul-3 conc: 2 function: 2 mass: 514.35>79.93



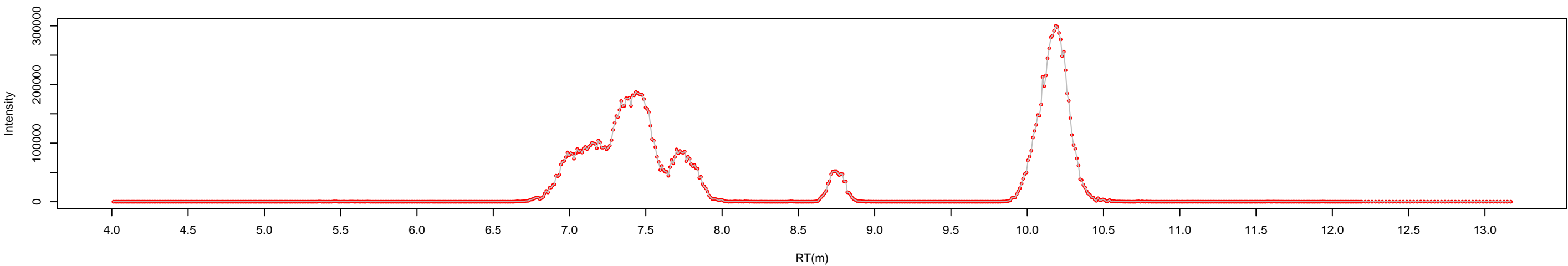
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 10.4 BLine: yes



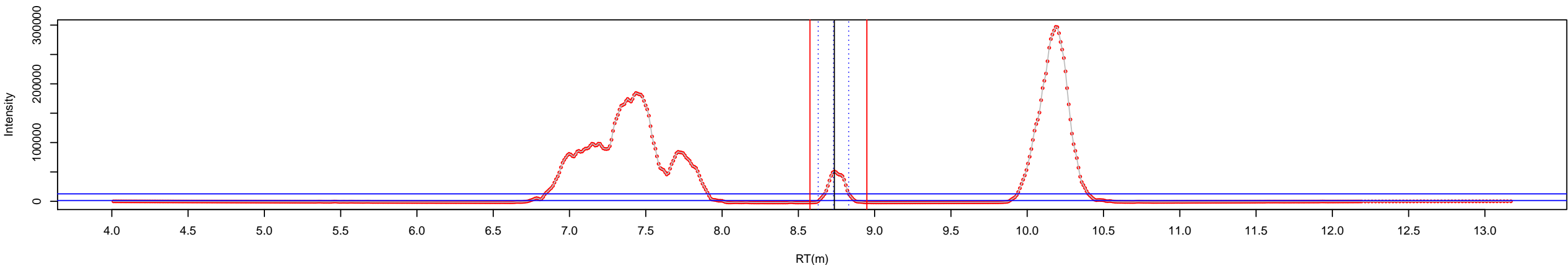
peak area | window size:3 BLine: yes



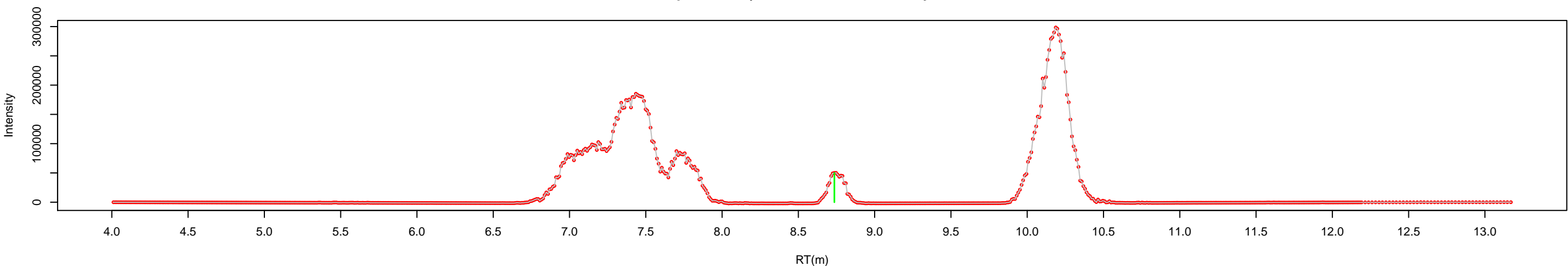
raw chromatogram | batch: 3 sample: std11-10ul-3 conc: 5 function: 2 mass: 514.35>79.93



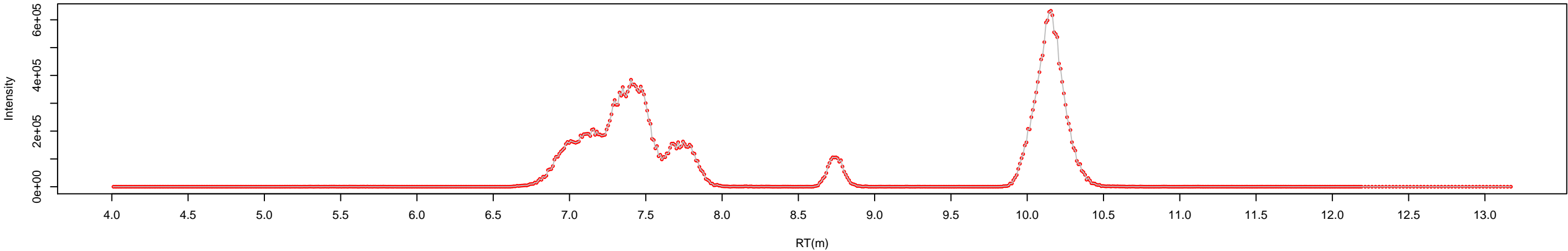
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 1270 BLine: yes



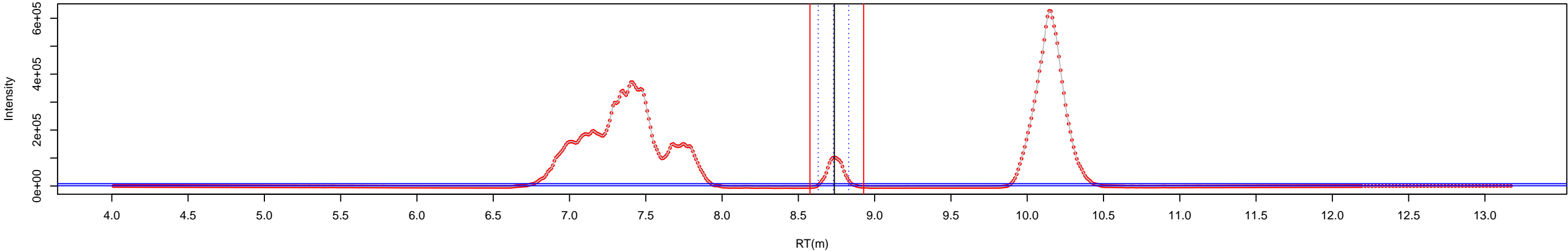
peak area | window size:3 BLine: yes



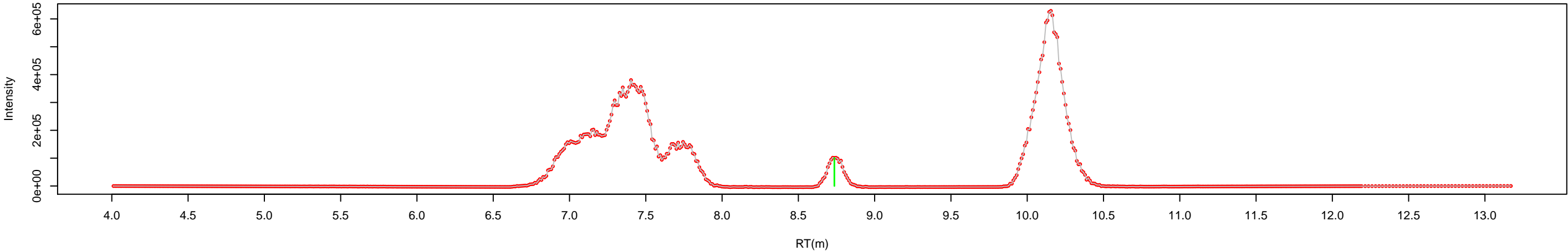
raw chromatogram | batch: 3 sample: std12-10ul-3 conc: 10 function: 2 mass: 514.35>79.93



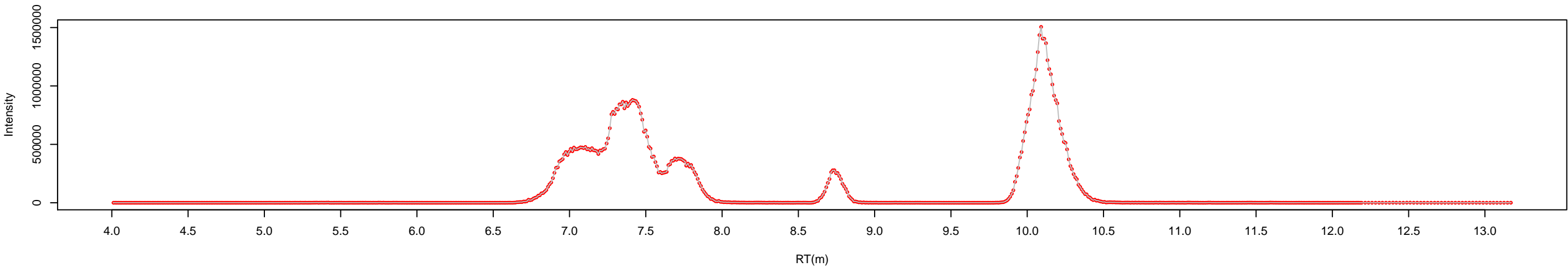
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 837 BLine: yes



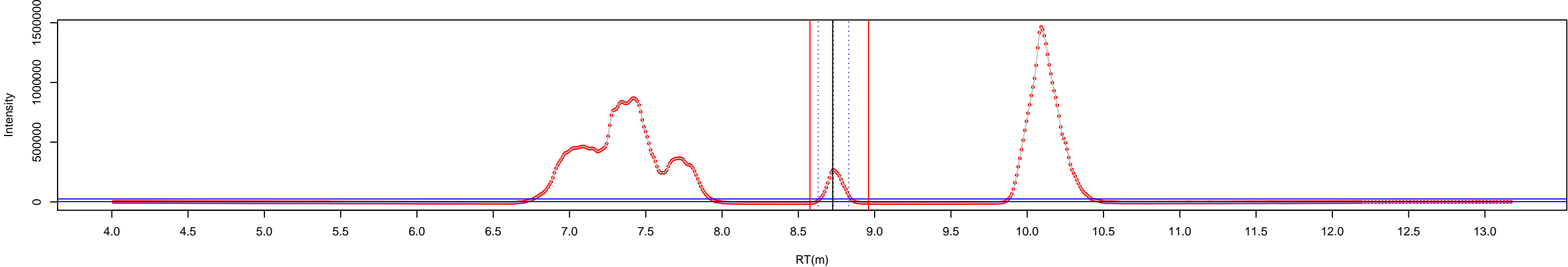
peak area | window size:3 BLine: yes



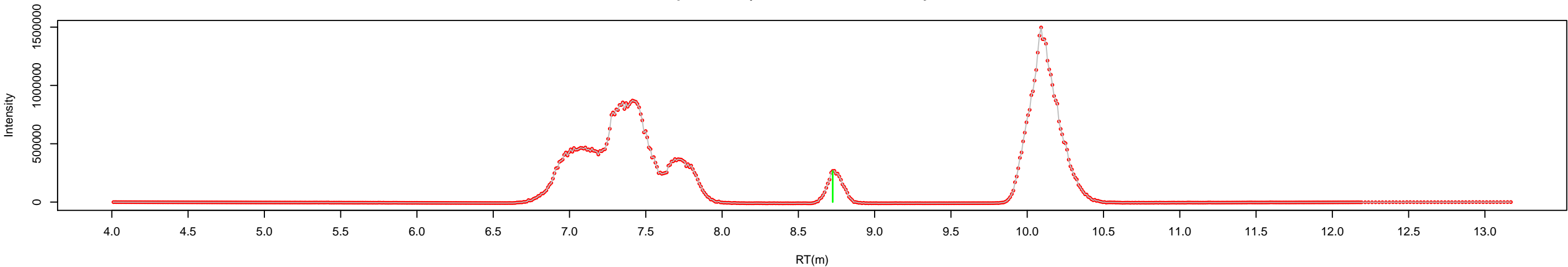
raw chromatogram | batch: 3 sample: std13-10ul-3 conc: 20 function: 2 mass: 514.35>79.93



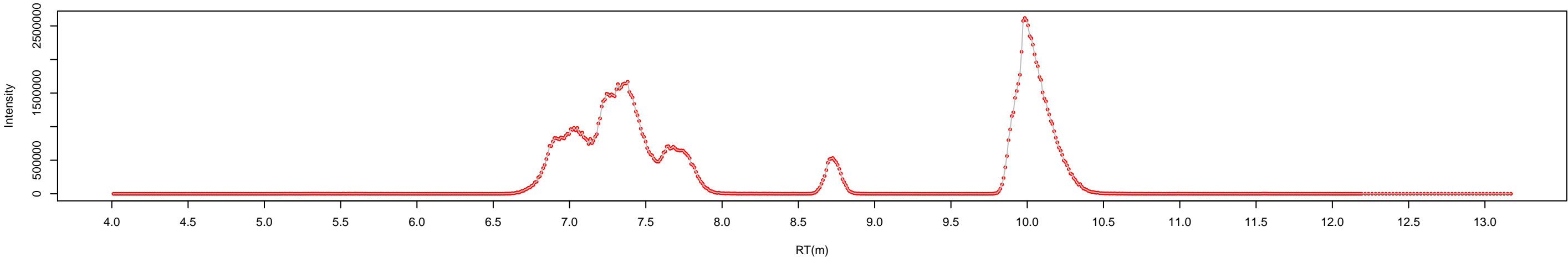
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 2460 BLine: yes



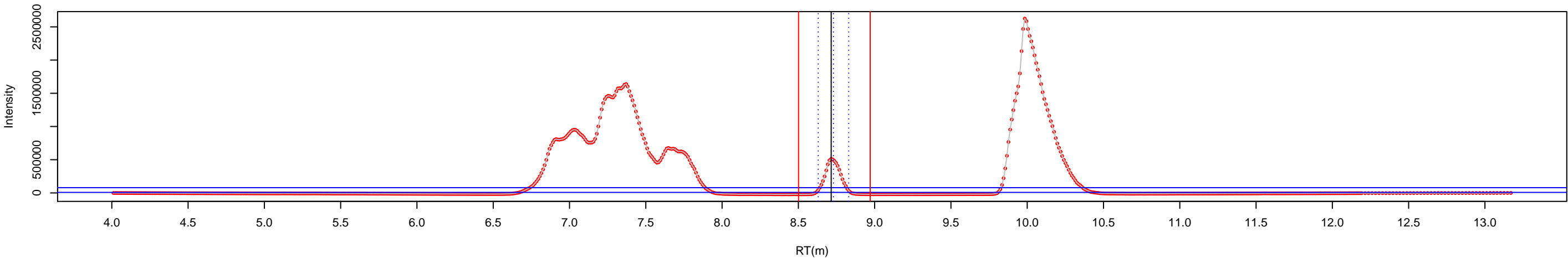
peak area | window size:3 BLine: yes



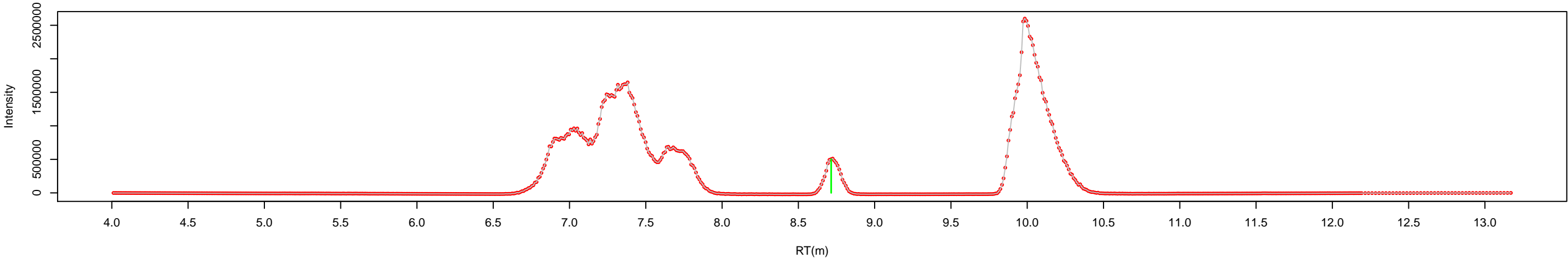
raw chromatogram | batch: 3   sample: std14-10ul-3   conc: 50   function: 2   mass: 514.35>79.93



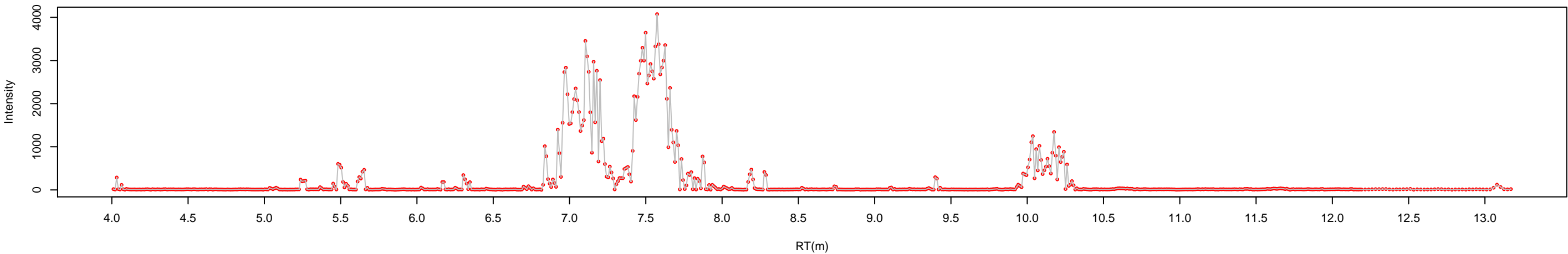
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 7880   BLine: yes



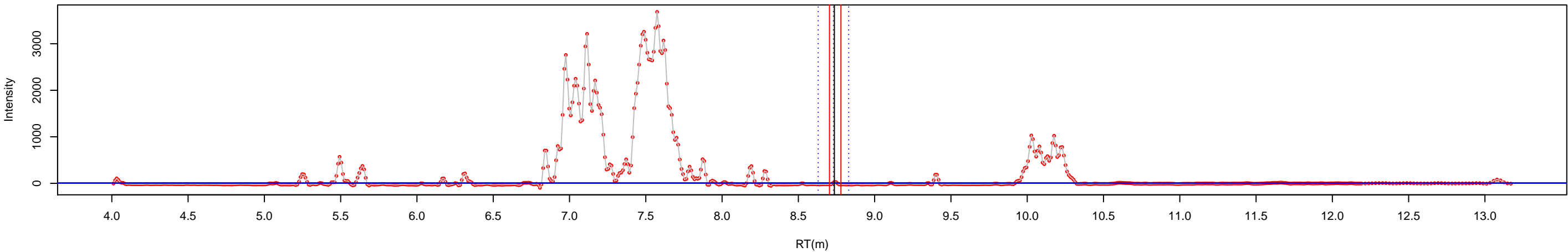
peak area | window size:3   BLine: yes



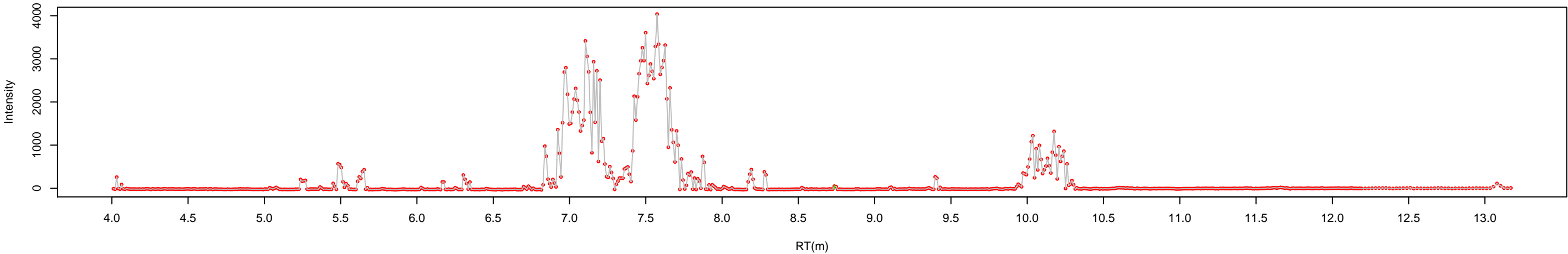
raw chromatogram | batch: 4 sample: cotor1 conc: NA function: 2 mass: 514.35>79.93



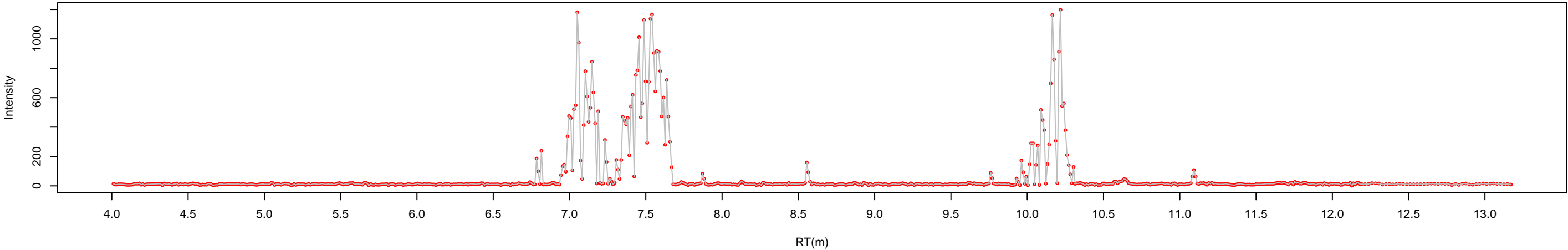
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 1.09 BLine: yes



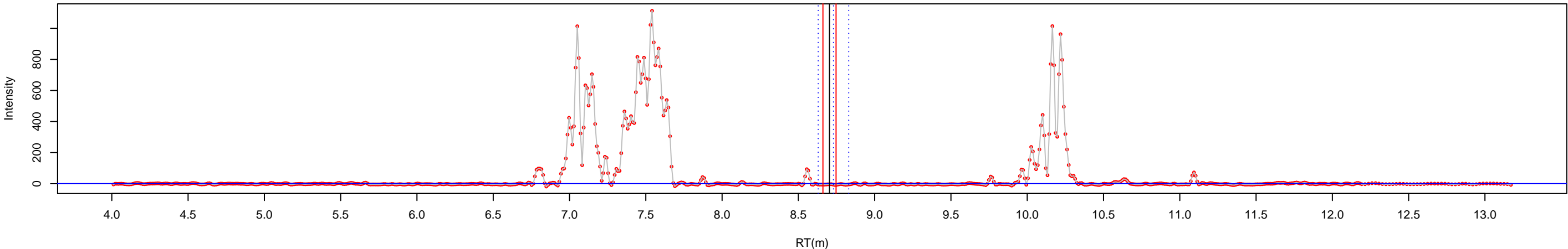
peak area | window size:3 BLine: yes



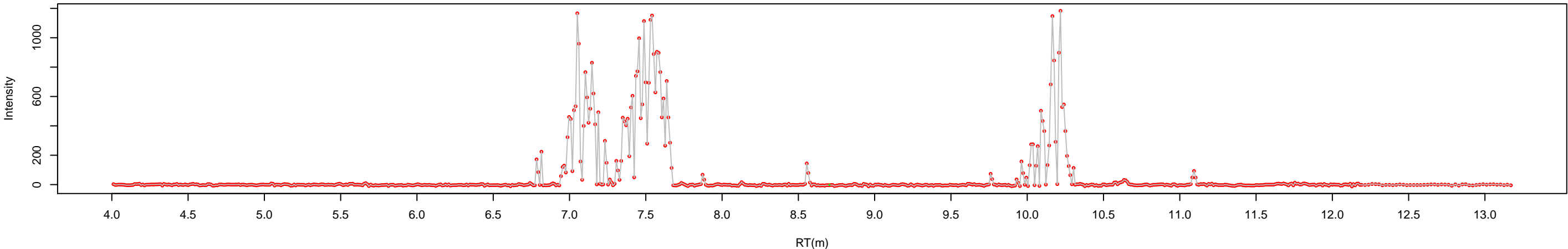
raw chromatogram | batch: 4   sample: cotor2   conc: NA   function: 2   mass: 514.35>79.93



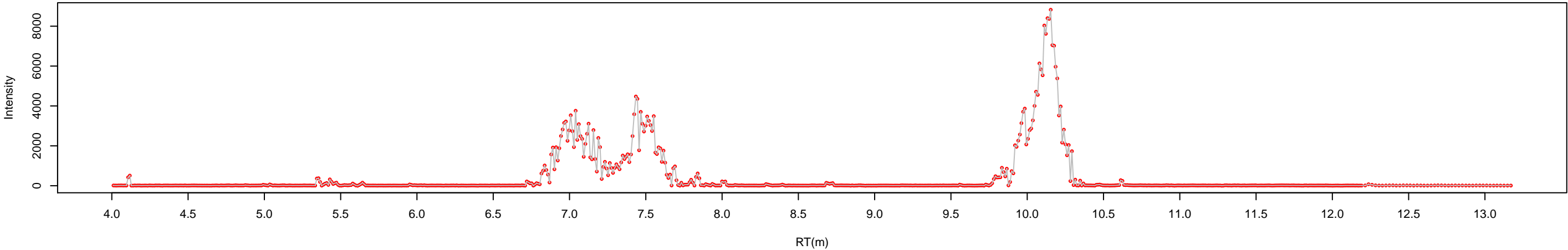
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 0.0532   BLine: yes



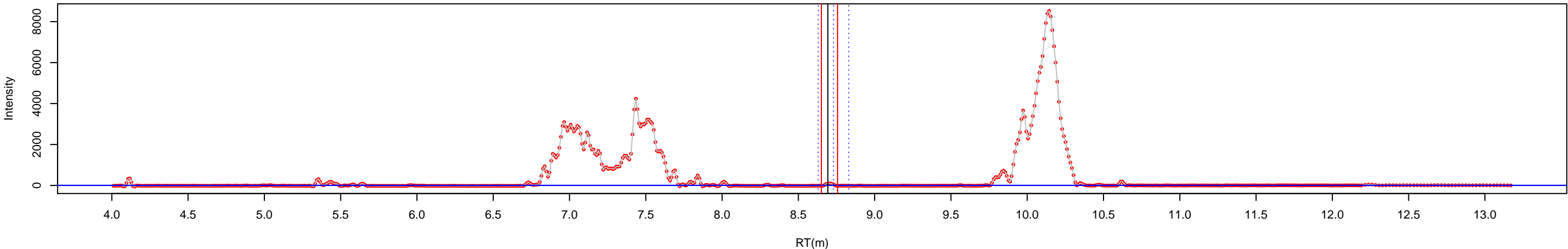
peak area | window size:3   BLine: yes



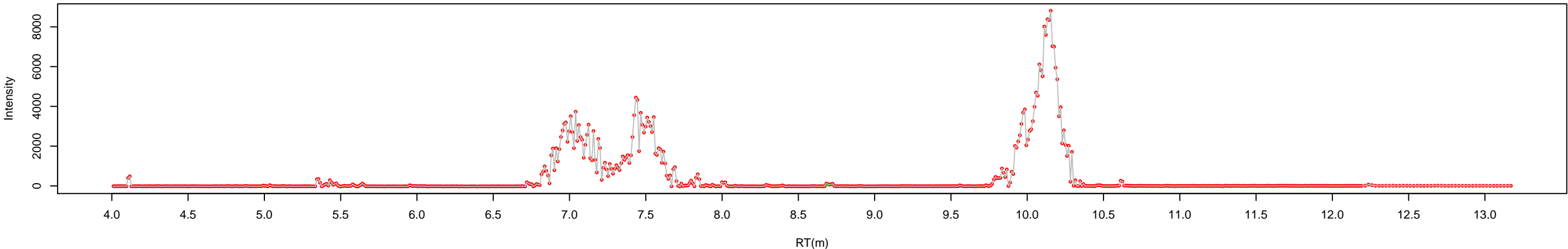
raw chromatogram | batch: 4   sample: cotor3   conc: NA   function: 2   mass: 514.35>79.93



peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 0.843   BLine: yes

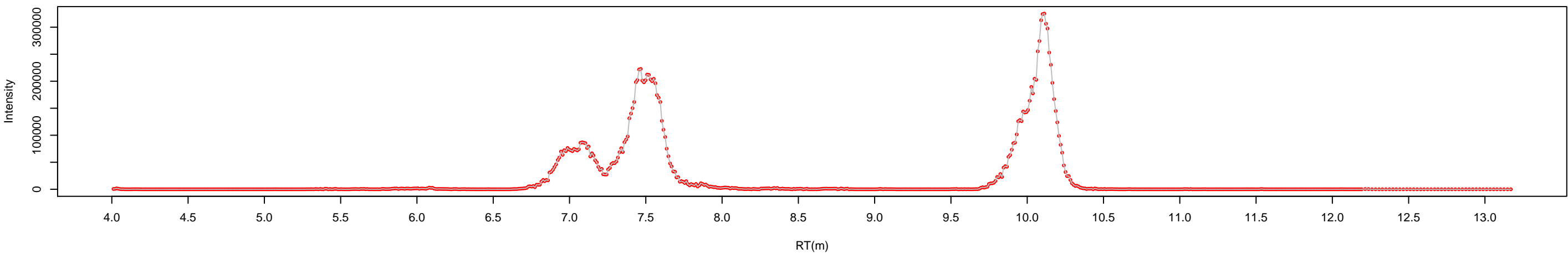


peak area | window size:3   BLine: yes

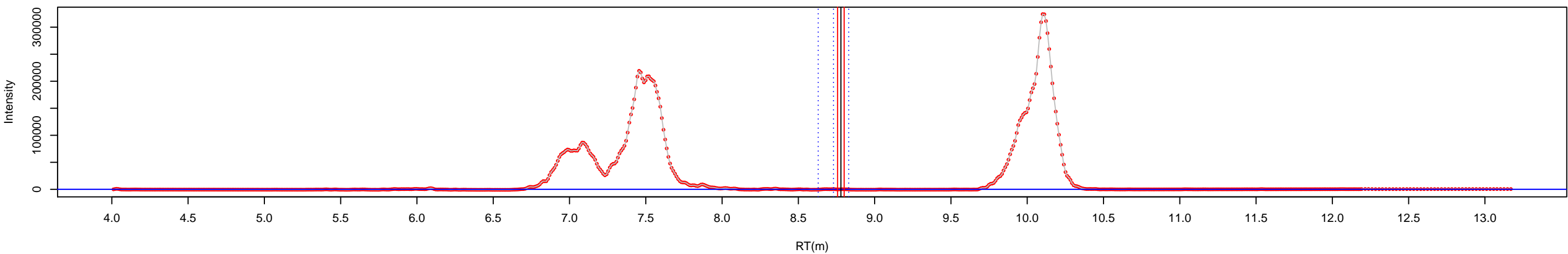




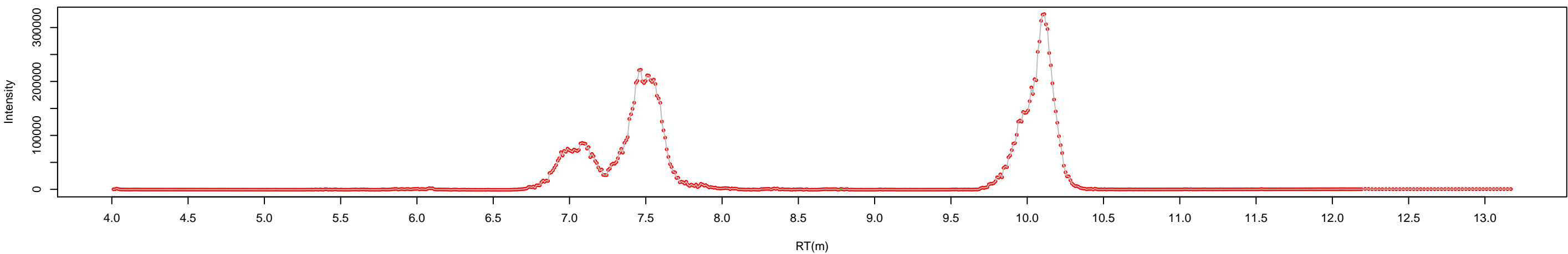
raw chromatogram | batch: 4   sample: cotor4   conc: NA   function: 2   mass: 514.35>79.93



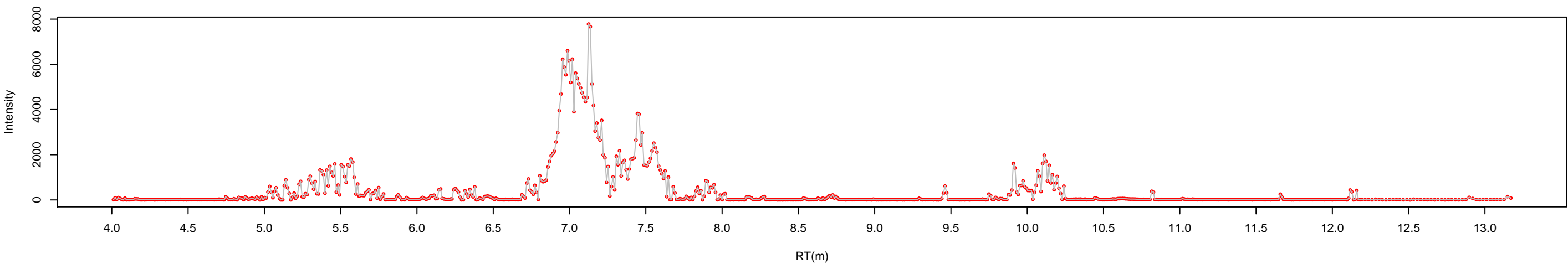
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 19   BLine: yes



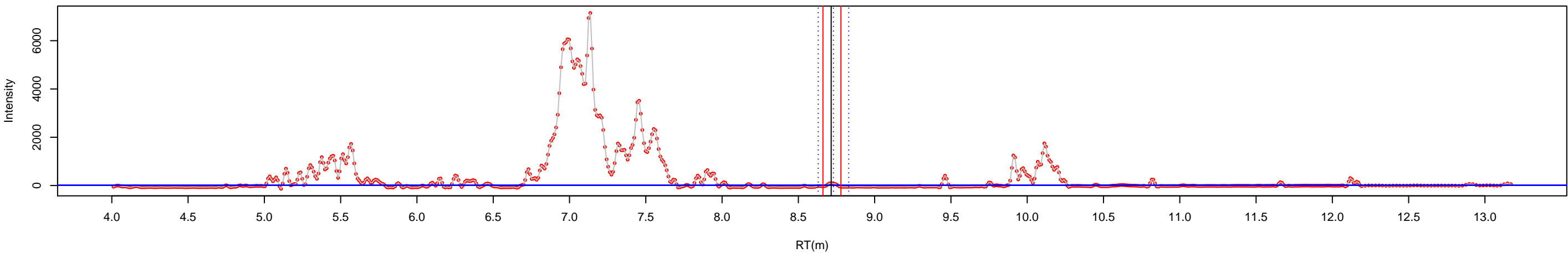
peak area | window size:3   BLine: yes



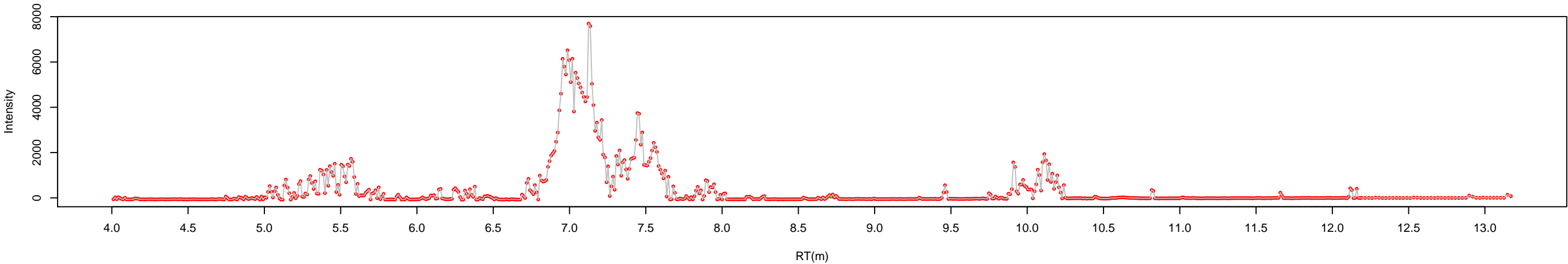
raw chromatogram | batch: 4 sample: cotor5 conc: NA function: 2 mass: 514.35>79.93



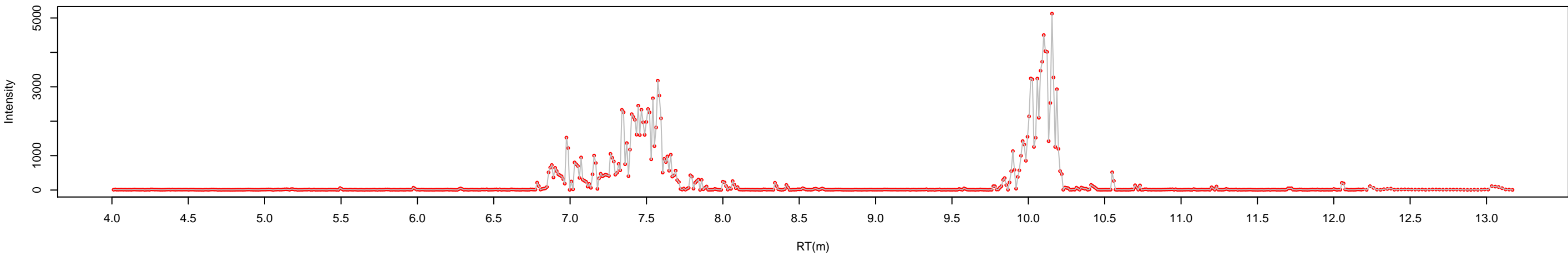
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 2.14 BLine: yes



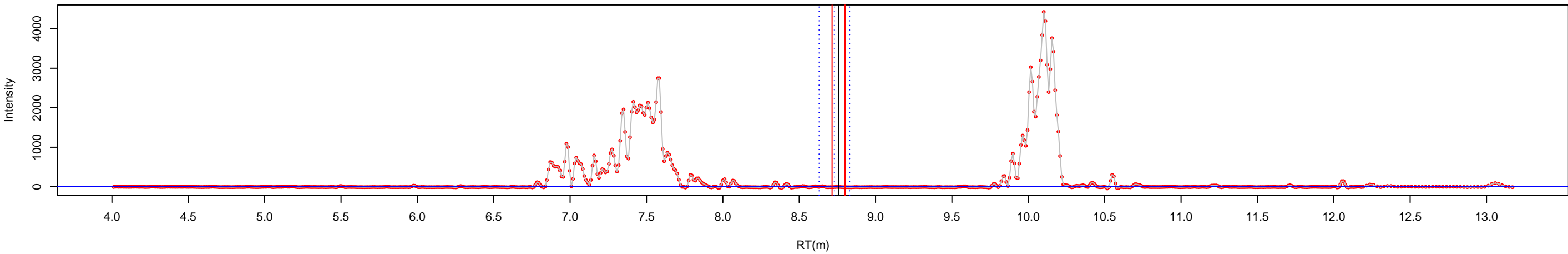
peak area | window size:3 BLine: yes



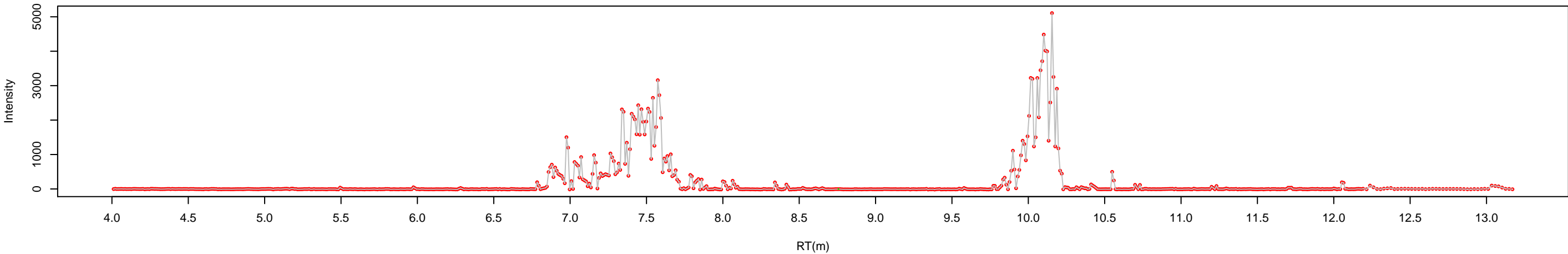
raw chromatogram | batch: 4 sample: cotor15 conc: NA function: 2 mass: 514.35>79.93



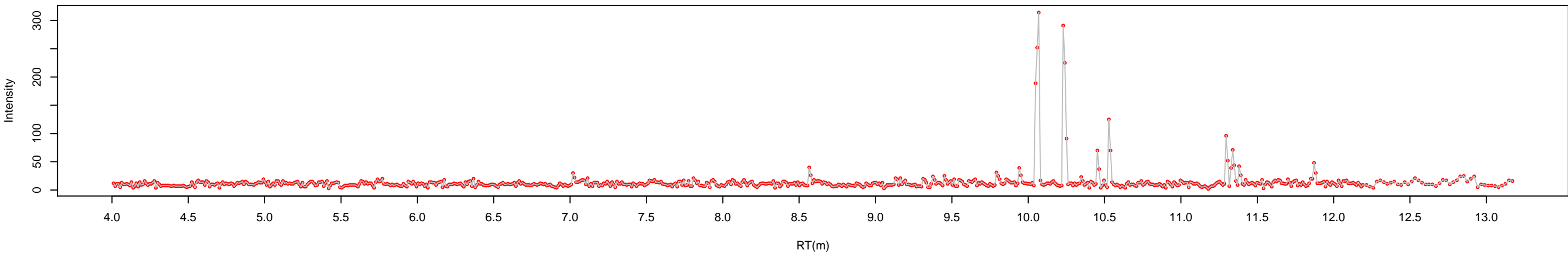
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.303 BLine: yes



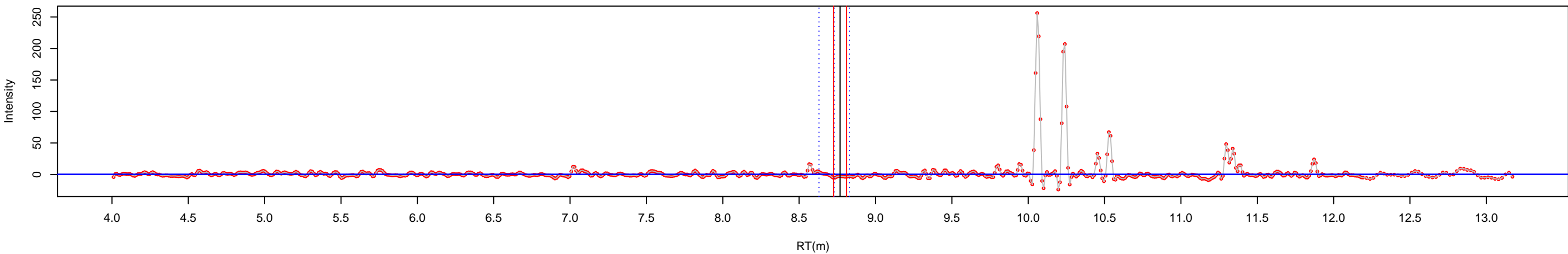
peak area | window size:3 BLine: yes



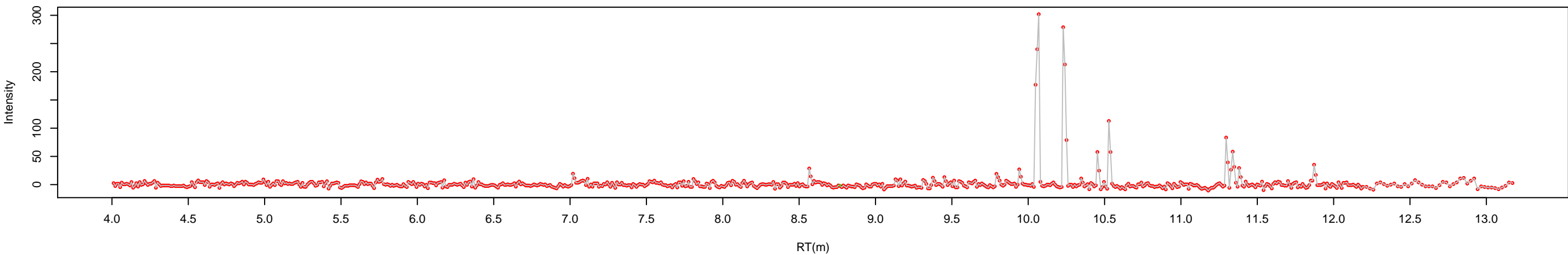
raw chromatogram | batch: 4 sample: std0-10ul-4 conc: 0.001 function: 2 mass: 514.35>79.93



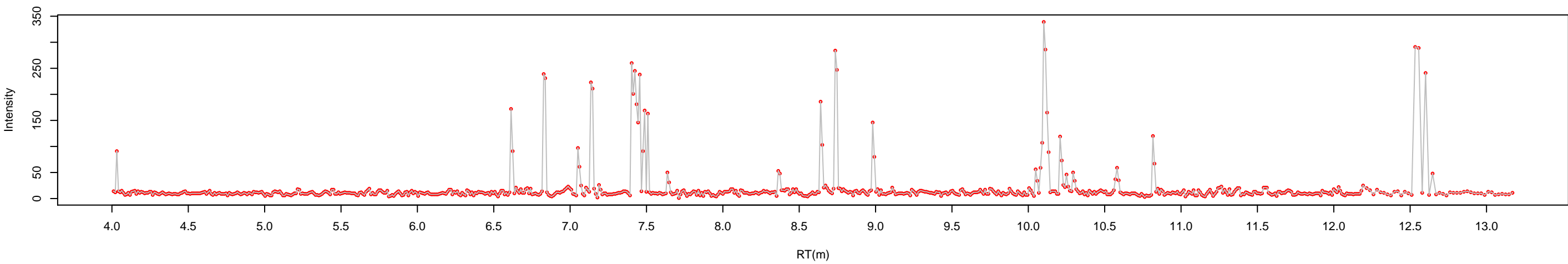
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.0574 BLine: yes



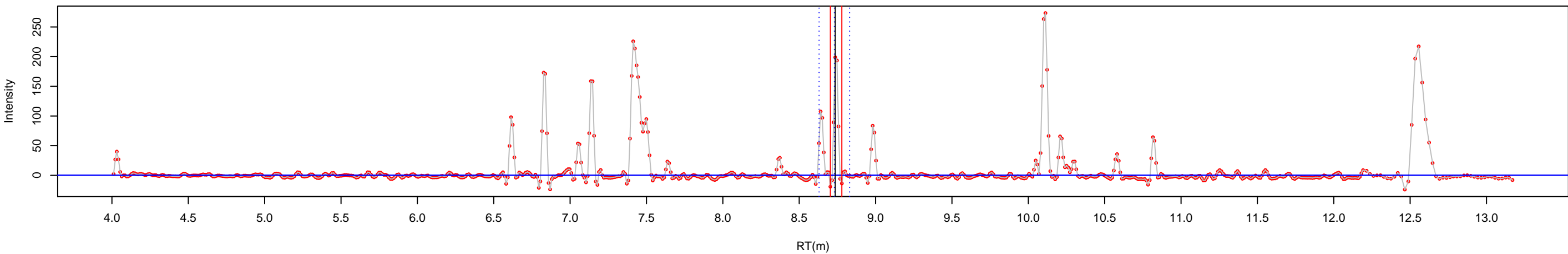
peak area | window size:3 BLine: yes



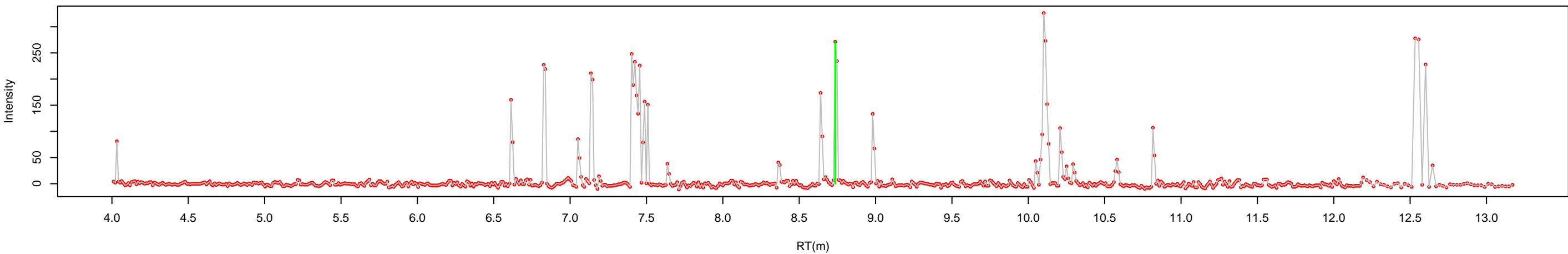
raw chromatogram | batch: 4 sample: std1-10ul-4 conc: 0.002 function: 2 mass: 514.35>79.93



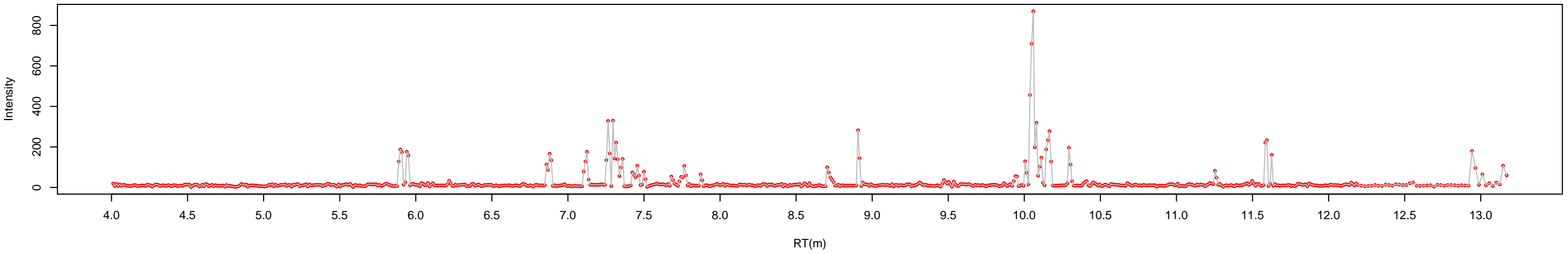
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.0371 BLine: yes



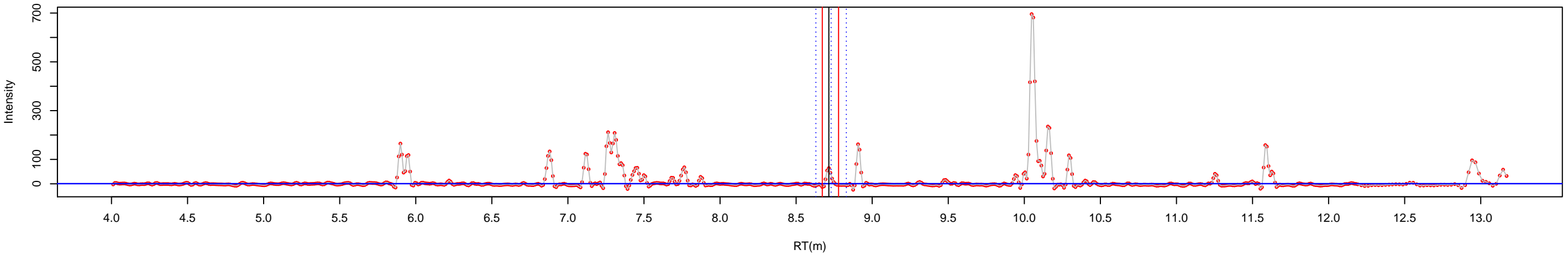
peak area | window size:3 BLine: yes



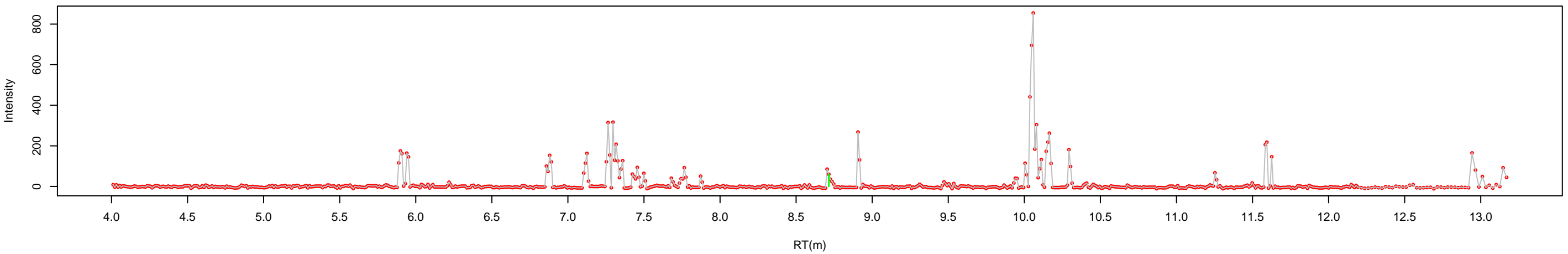
raw chromatogram | batch: 4   sample: std2-10ul-4   conc: 0.005   function: 2   mass: 514.35>79.93



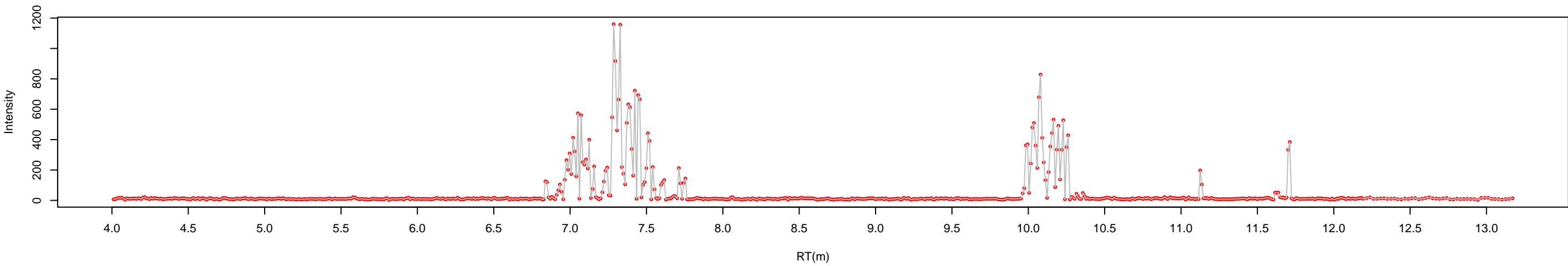
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 0.122   BLine: yes



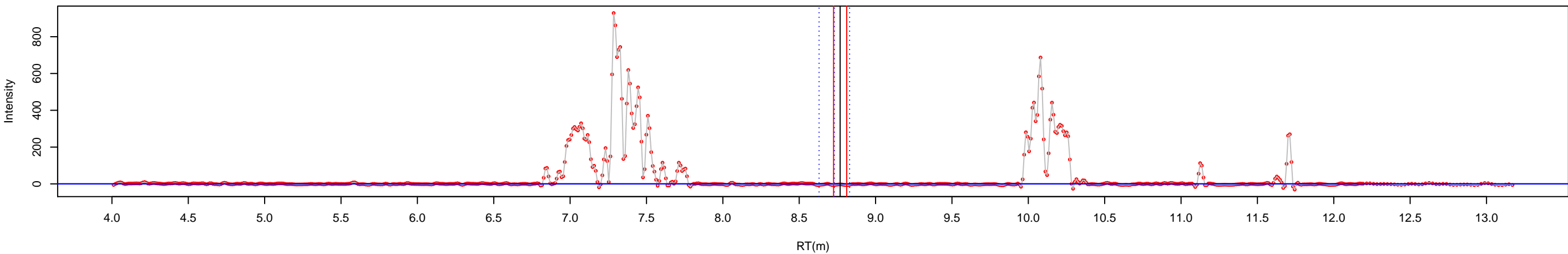
peak area | window size:3   BLine: yes



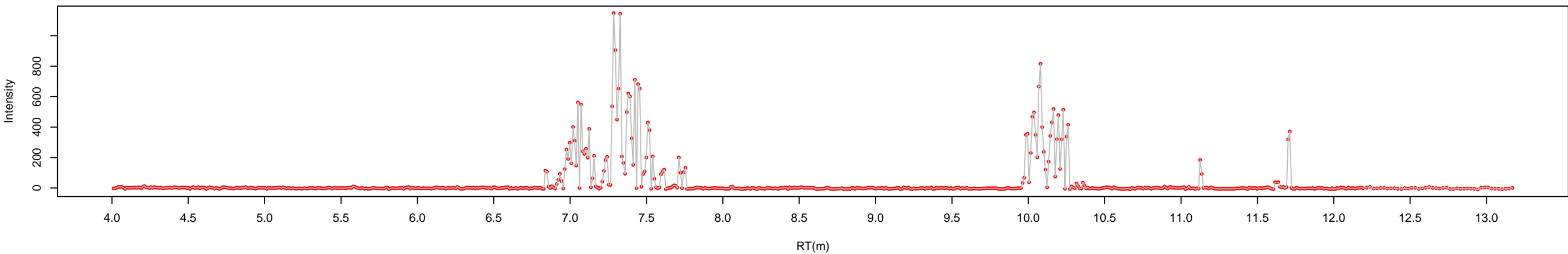
raw chromatogram | batch: 4 sample: std3-10ul-4 conc: 0.01 function: 2 mass: 514.35>79.93



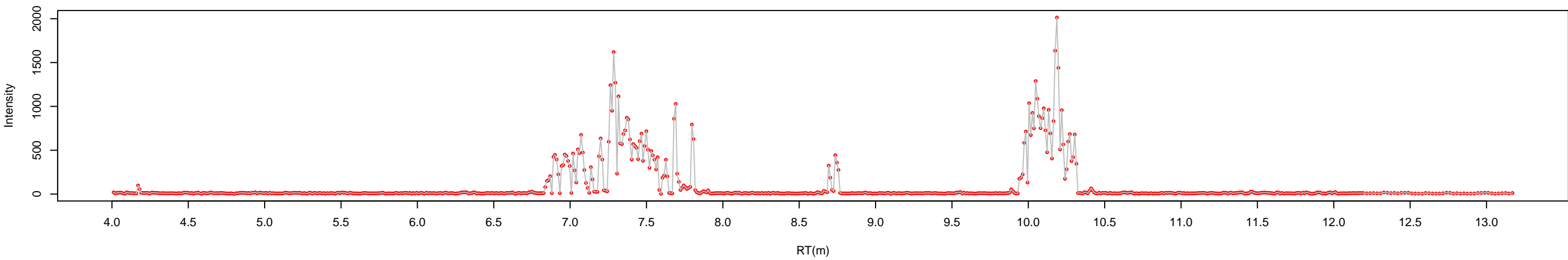
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.0414 BLine: yes



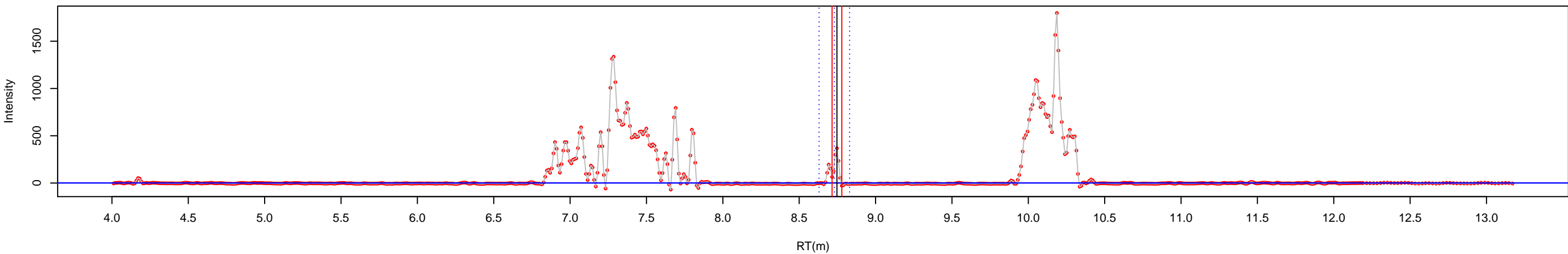
peak area | window size:3 BLine: yes



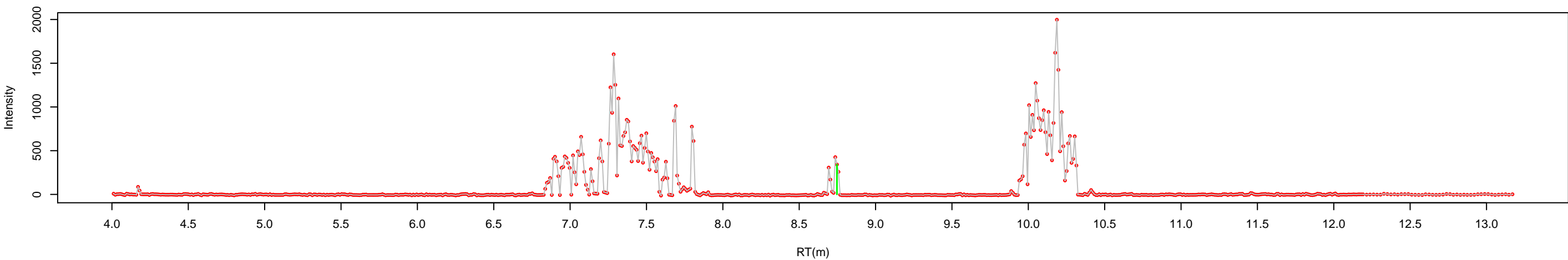
raw chromatogram | batch: 4 sample: std4-10ul-4 conc: 0.02 function: 2 mass: 514.35>79.93



peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.155 BLine: yes

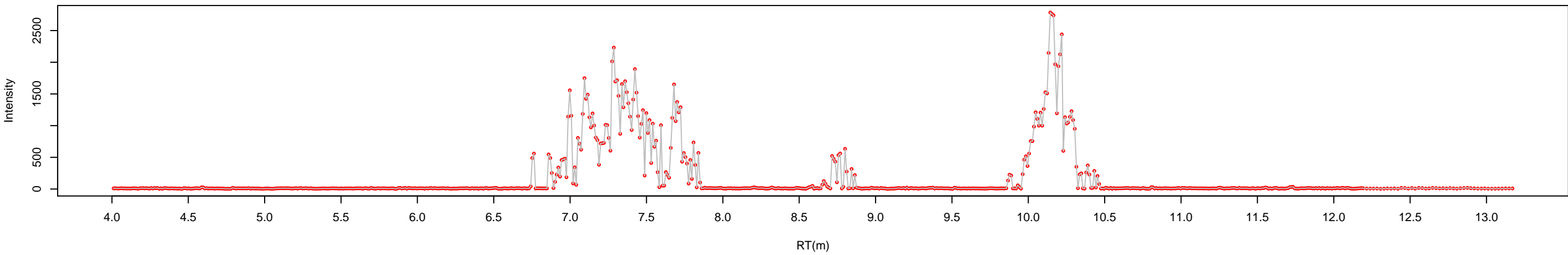


peak area | window size:3 BLine: yes

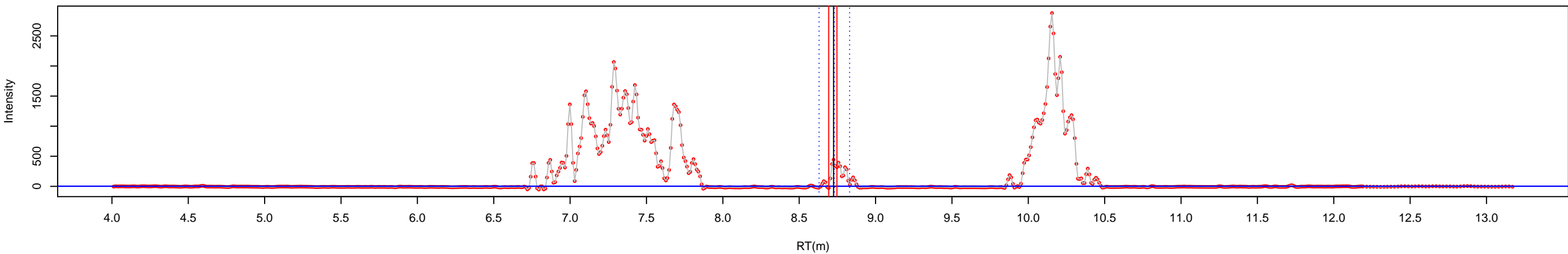




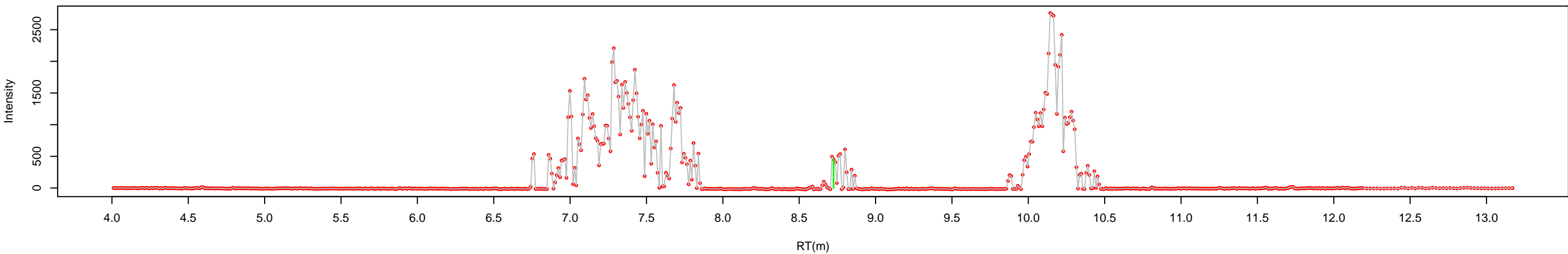
raw chromatogram | batch: 4 sample: std5-10ul-4 conc: 0.05 function: 2 mass: 514.35>79.93



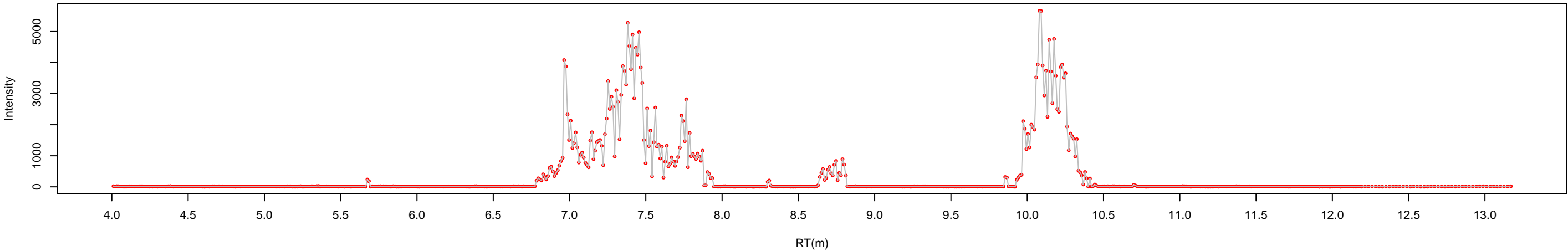
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 0.425 BLine: yes



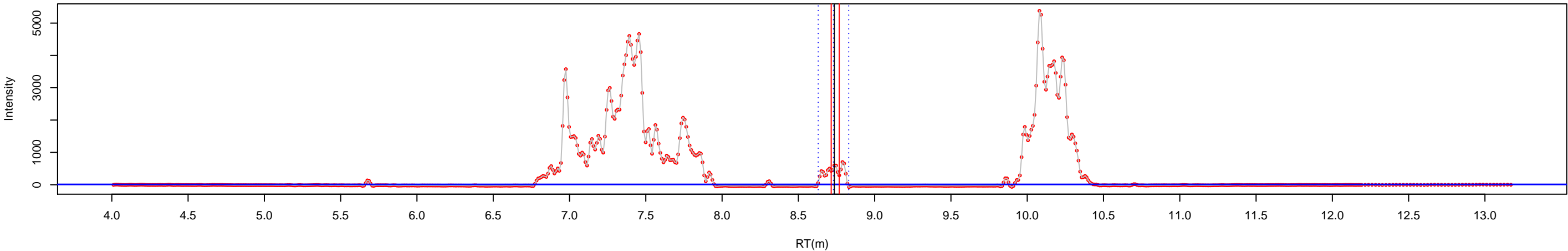
peak area | window size:3 BLine: yes



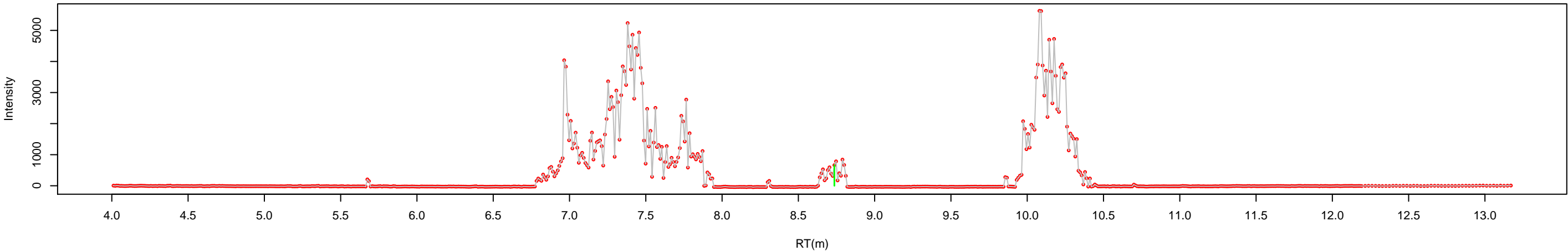
raw chromatogram | batch: 4   sample: std6-10ul-4   conc: 0.1   function: 2   mass: 514.35>79.93



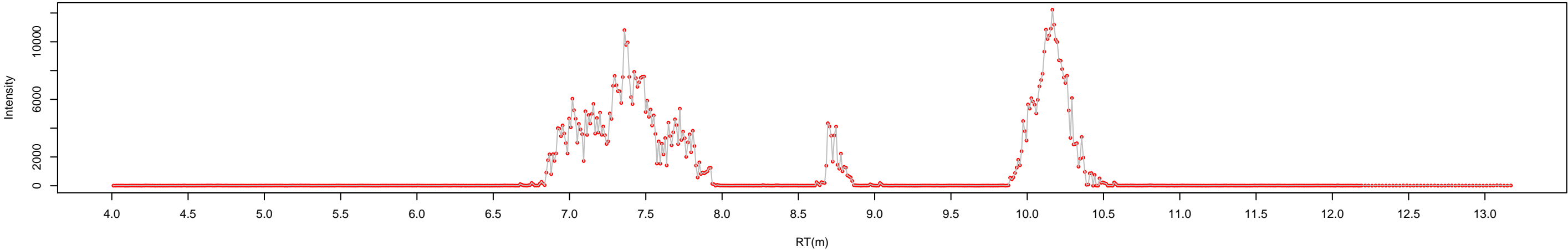
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 1.73   BLine: yes



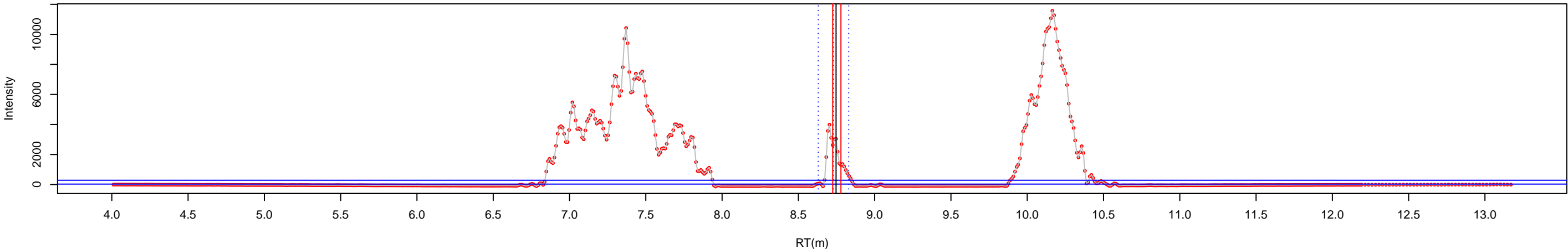
peak area | window size:3   BLine: yes



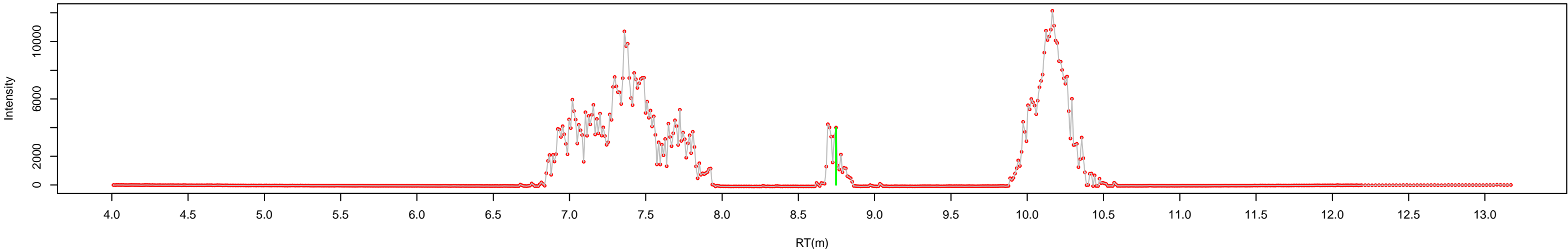
raw chromatogram | batch: 4   sample: std7-10ul-4   conc: 0.2   function: 2   mass: 514.35>79.93



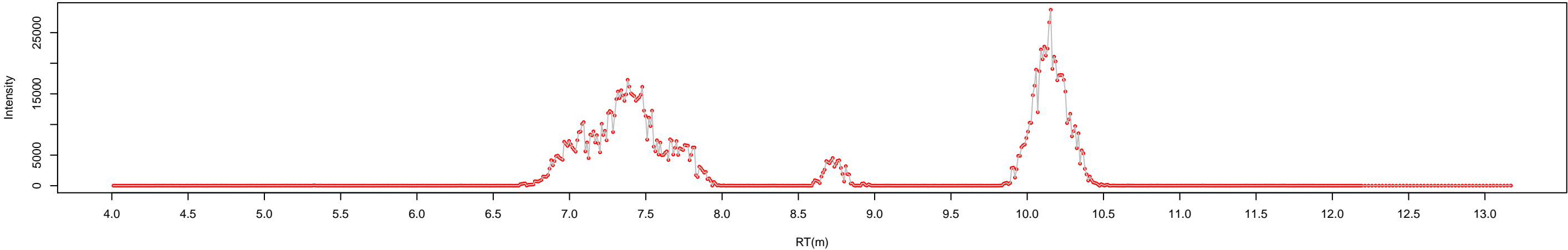
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 28.5   BLine: yes



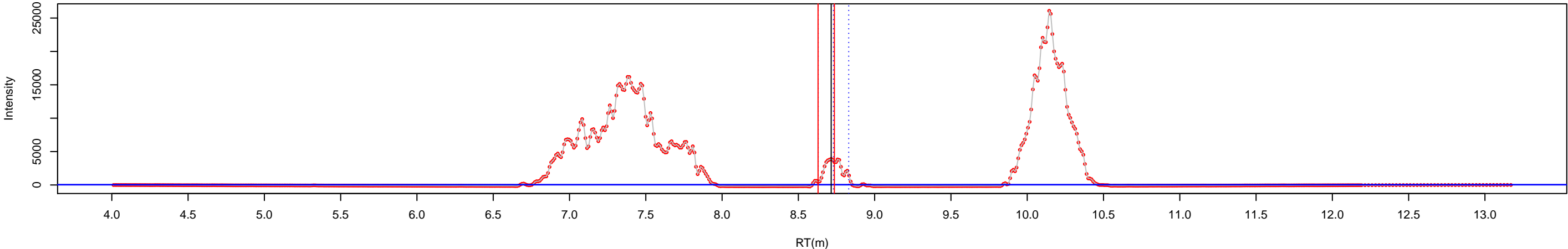
peak area | window size:3   BLine: yes



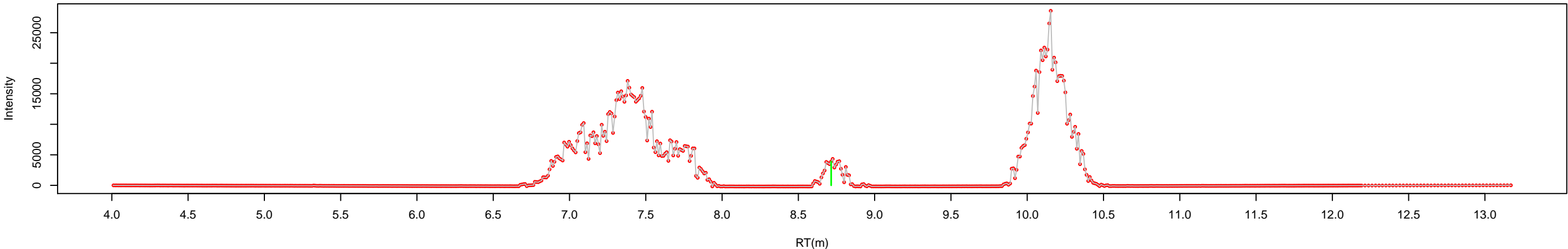
raw chromatogram | batch: 4   sample: std8-10ul-4   conc: 0.5   function: 2   mass: 514.35>79.93



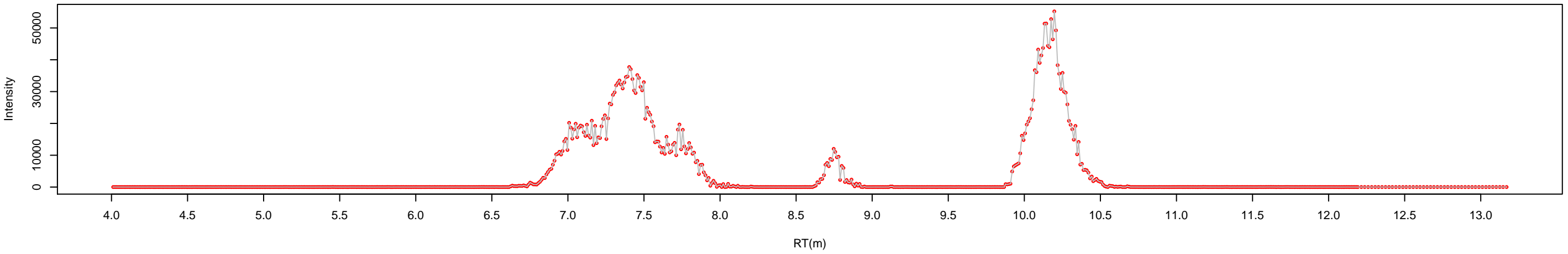
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 7.11   BLine: yes



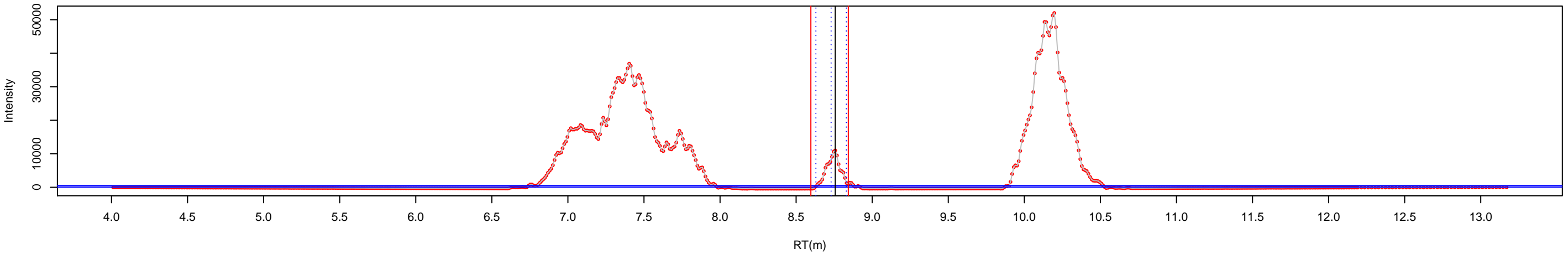
peak area | window size:3   BLine: yes



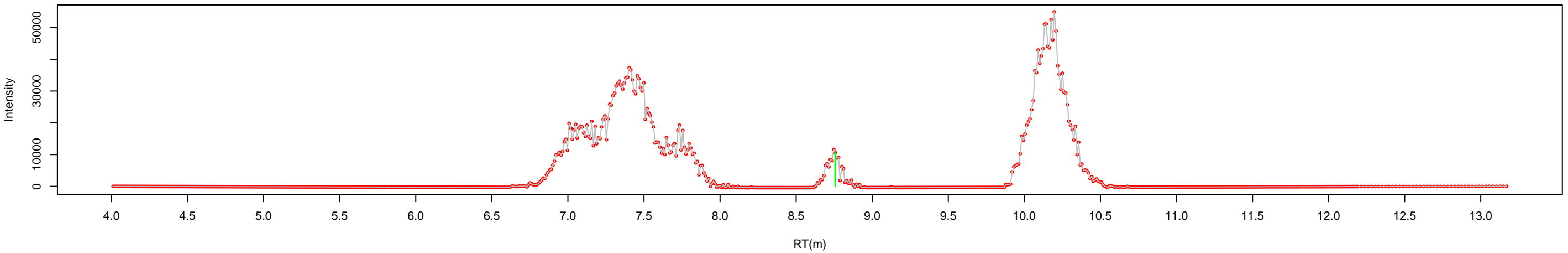
raw chromatogram | batch: 4   sample: std9-10ul-4   conc: 1   function: 2   mass: 514.35>79.93



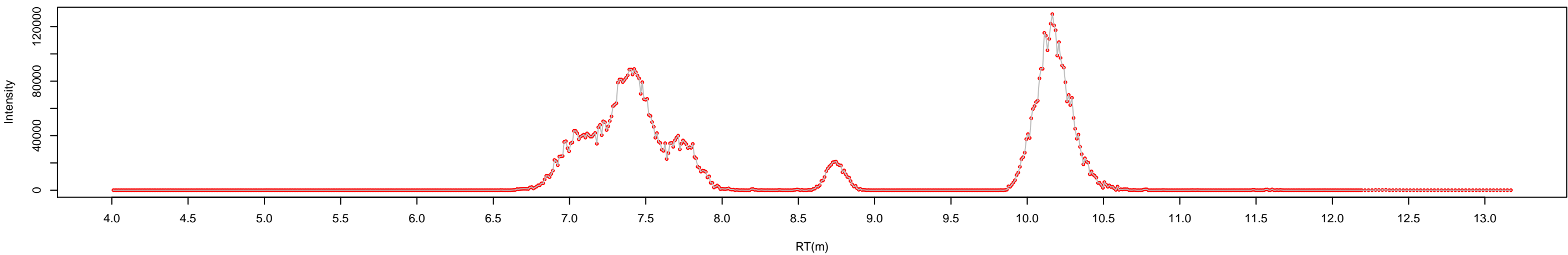
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 49.5   BLine: yes



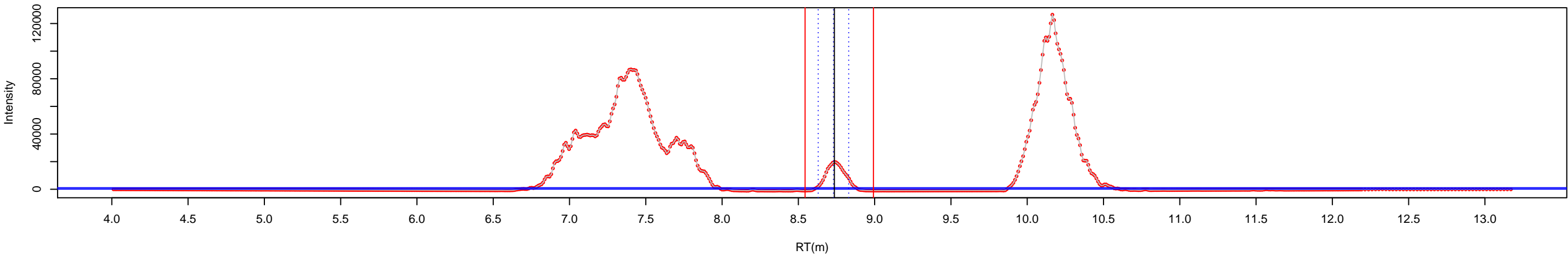
peak area | window size:3   BLine: yes



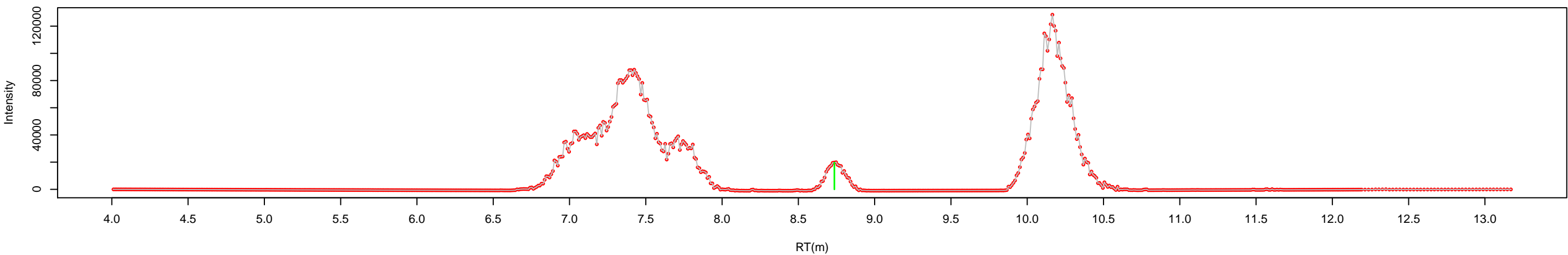
raw chromatogram | batch: 4   sample: std10-10ul-4   conc: 2   function: 2   mass: 514.35>79.93



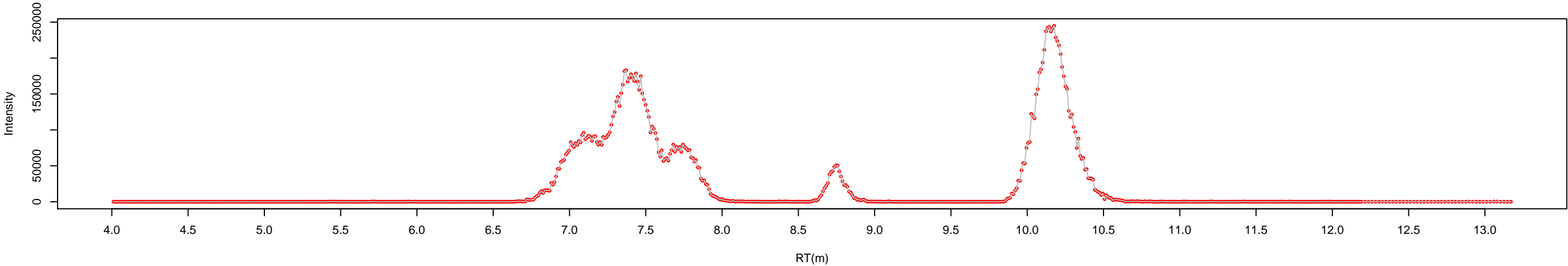
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 110   BLine: yes



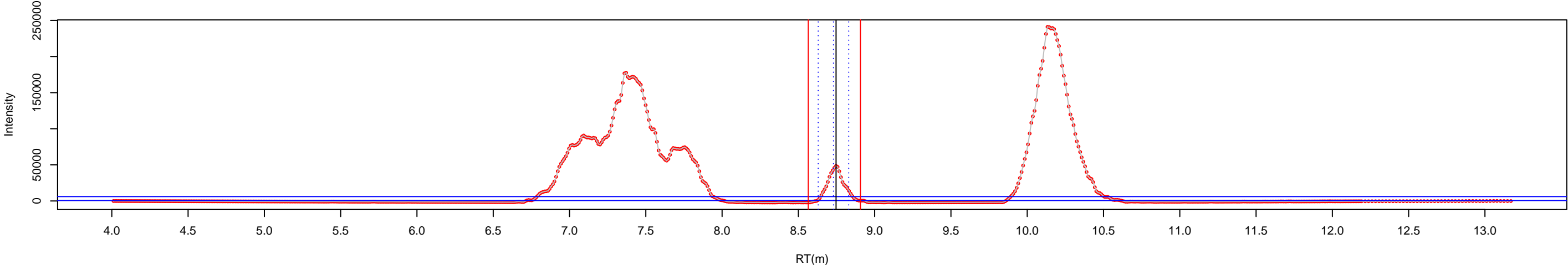
peak area | window size:3   BLine: yes



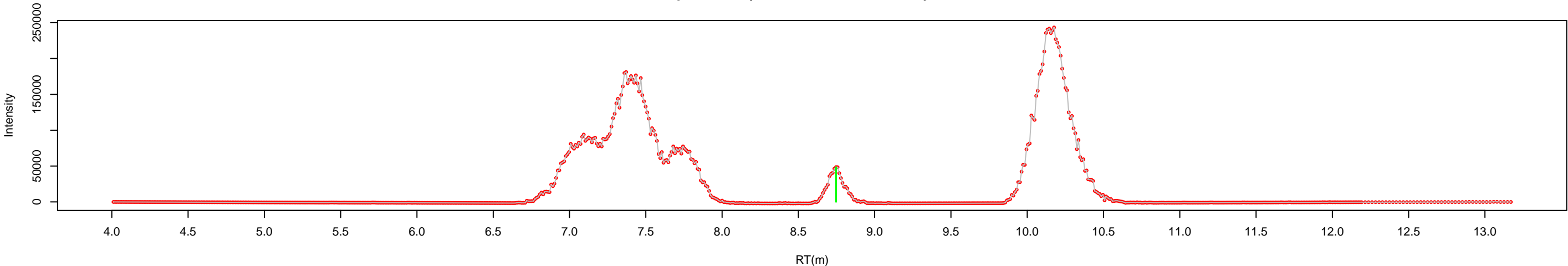
raw chromatogram | batch: 4 sample: std11-10ul-4 conc: 5 function: 2 mass: 514.35>79.93



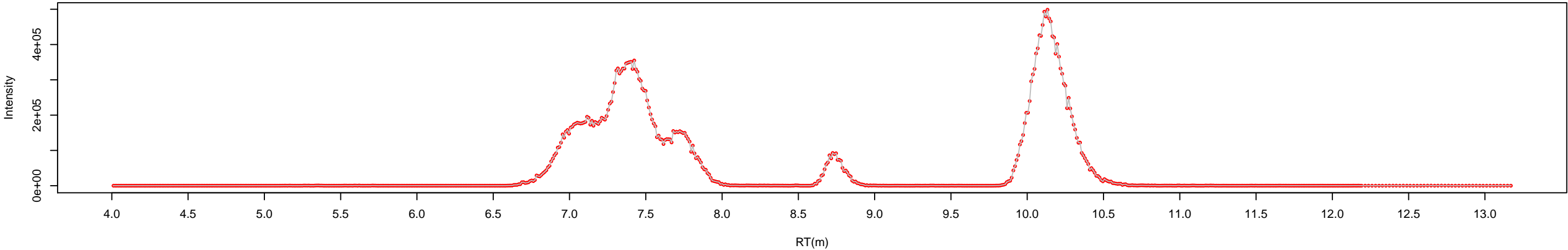
peak picking | window size: 7 iteration: 5 lp: 1 rp: 1 snr: 10 peak location: Nearest noise: 607 BLine: yes



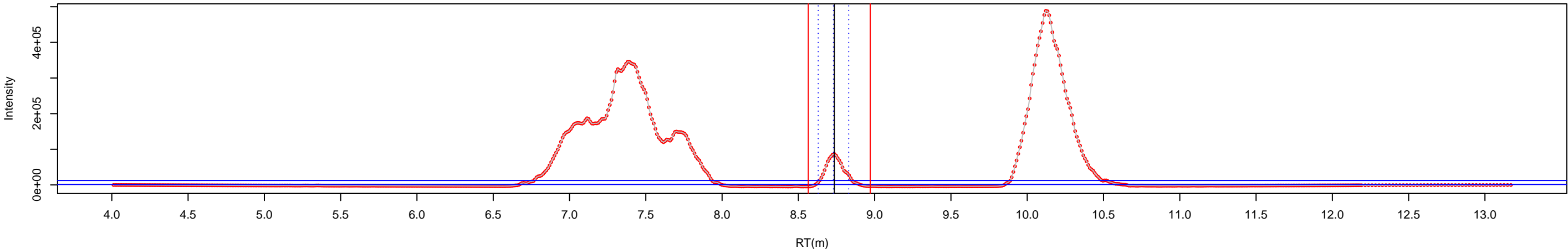
peak area | window size:3 BLine: yes



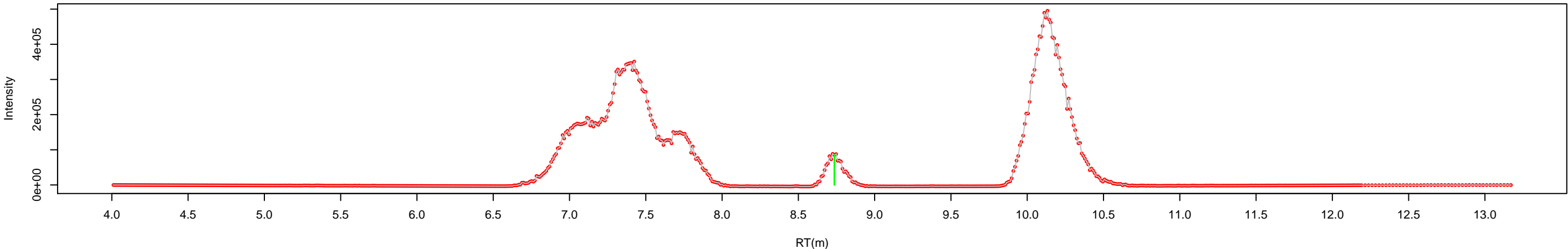
raw chromatogram | batch: 4   sample: std12-10ul-4   conc: 10   function: 2   mass: 514.35>79.93



peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 1250   BLine: yes

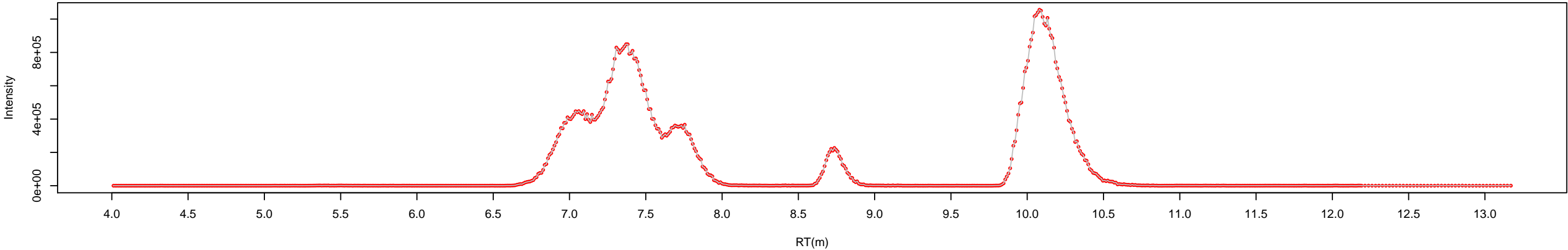


peak area | window size:3   BLine: yes

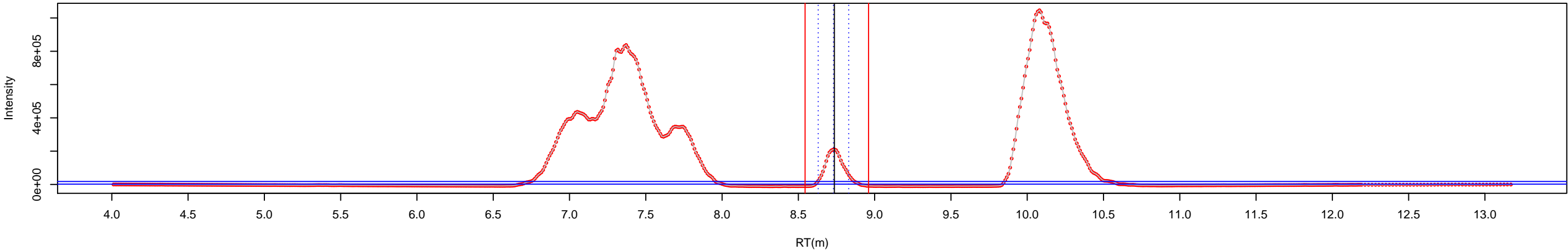




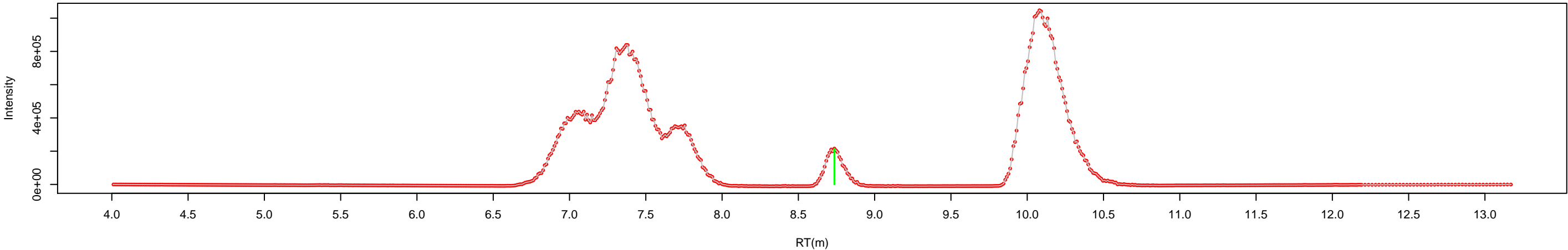
raw chromatogram | batch: 4   sample: std13-10ul-4   conc: 20   function: 2   mass: 514.35>79.93



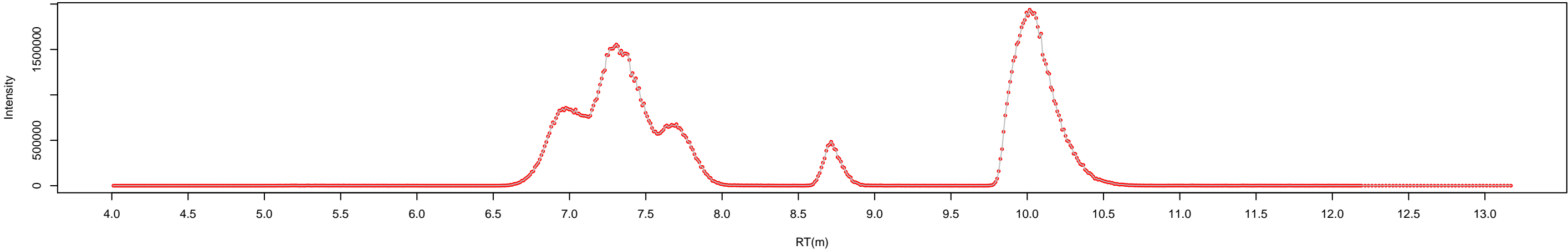
peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 1770   BLine: yes



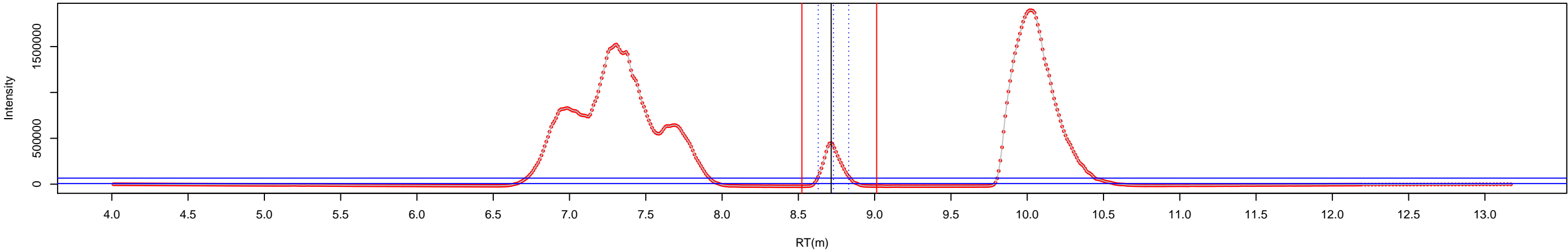
peak area | window size:3   BLine: yes



raw chromatogram | batch: 4   sample: std14-10ul-4   conc: 50   function: 2   mass: 514.35>79.93



peak picking | window size: 7   iteration: 5   lp: 1   rp: 1   snr: 10   peak location: Nearest   noise: 6560   BLine: yes



peak area | window size:3   BLine: yes

