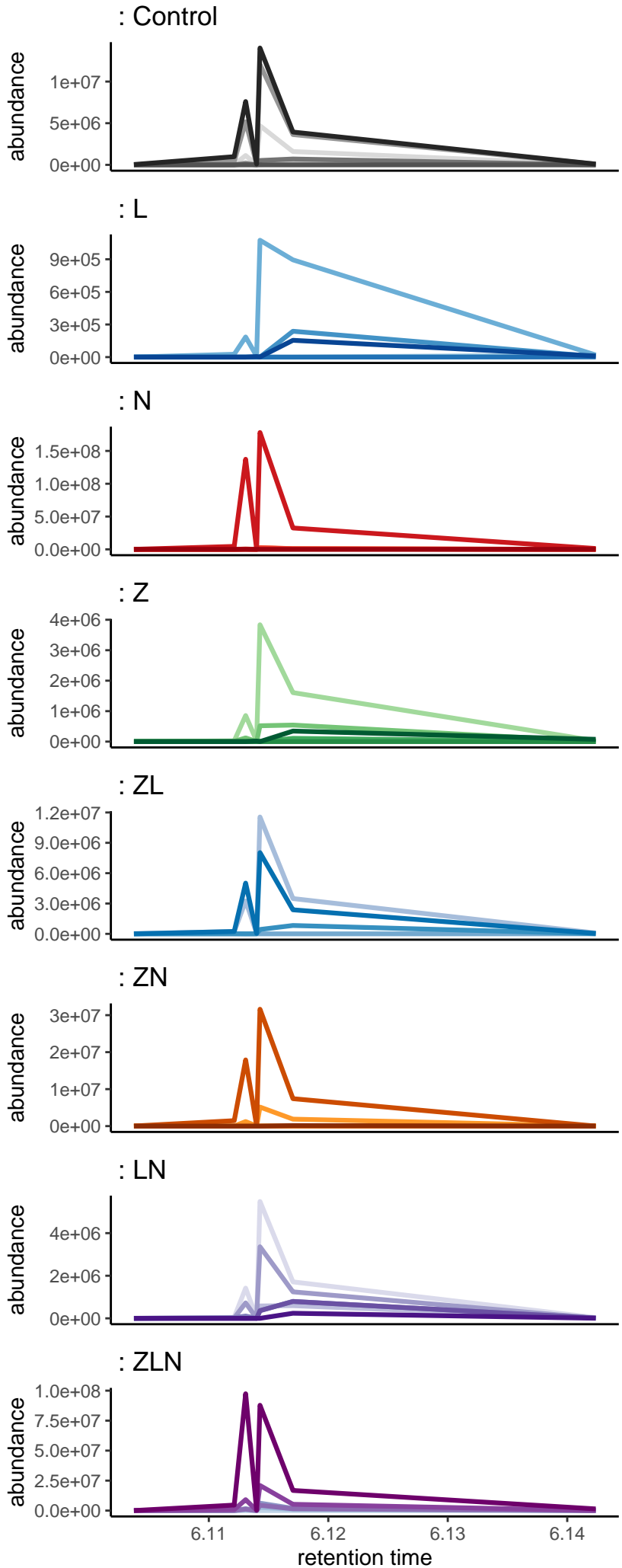
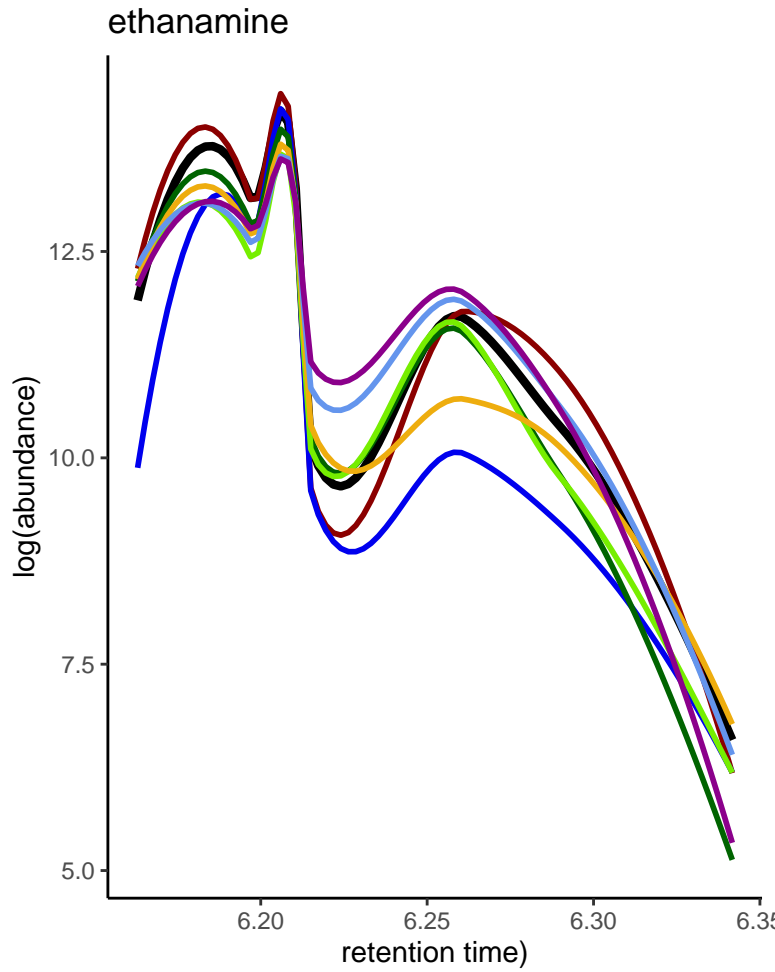


colour    c\_mean    ln\_mean    z\_mean    zln\_mean  
           l\_mean    n\_mean    zl\_mean    zn\_mean

X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
1	6.103733	0.26	0.42	0.28	0.00	0.84	0.54	0.38
2	6.112125	0.73	0.43	0.21	0.01	0.72	0.26	0.46
4	6.113083	0.94	0.65	0.18	0.00	0.68	0.27	0.75
6	6.114000	0.39	0.74	0.48	0.01	0.71	0.21	0.84
7	6.114283	0.63	0.58	0.11	0.00	0.64	0.14	0.36
9	6.117042	0.92	0.45	0.44	0.16	0.37	0.04	0.66
18	6.142383	0.78	0.37	0.34	0.04	0.24	0.62	0.72

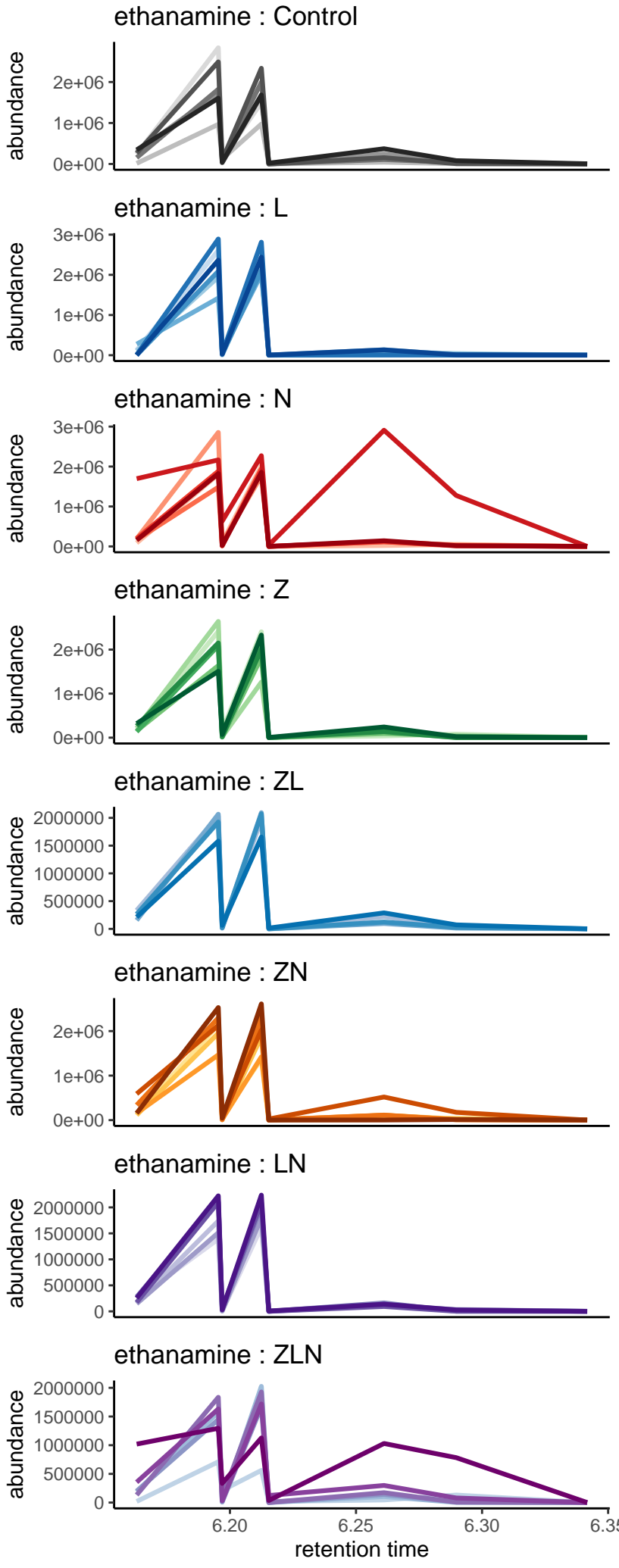


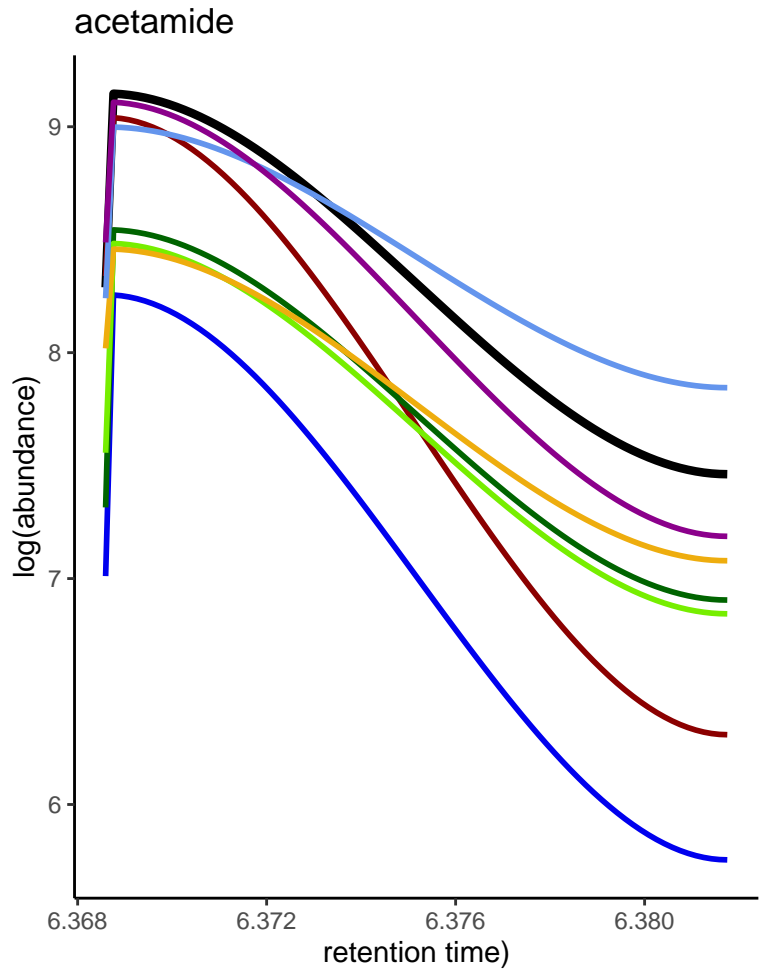


colour

c_mean	ln_mean	z_mean	zln_mean
l_mean	n_mean	zl_mean	zn_mean

ethanamine								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
21	6.163000	0.10	0.11	0.05	0.14	0.04	0.21	0.13
40	6.195392	0.37	0.18	0.20	0.04	0.54	0.06	0.78
42	6.196917	0.84	0.14	0.93	0.03	0.42	0.57	0.36
60	6.212467	0.19	0.96	0.24	0.03	0.29	0.03	0.84
63	6.215400	0.02	0.35	0.64	0.21	0.66	0.28	0.52
86	6.261108	0.61	0.79	0.45	0.02	0.10	0.11	0.73
88	6.289767	0.50	0.38	0.23	0.04	0.97	0.35	0.95
95	6.341700	0.23	0.62	0.82	0.97	0.43	0.19	0.04

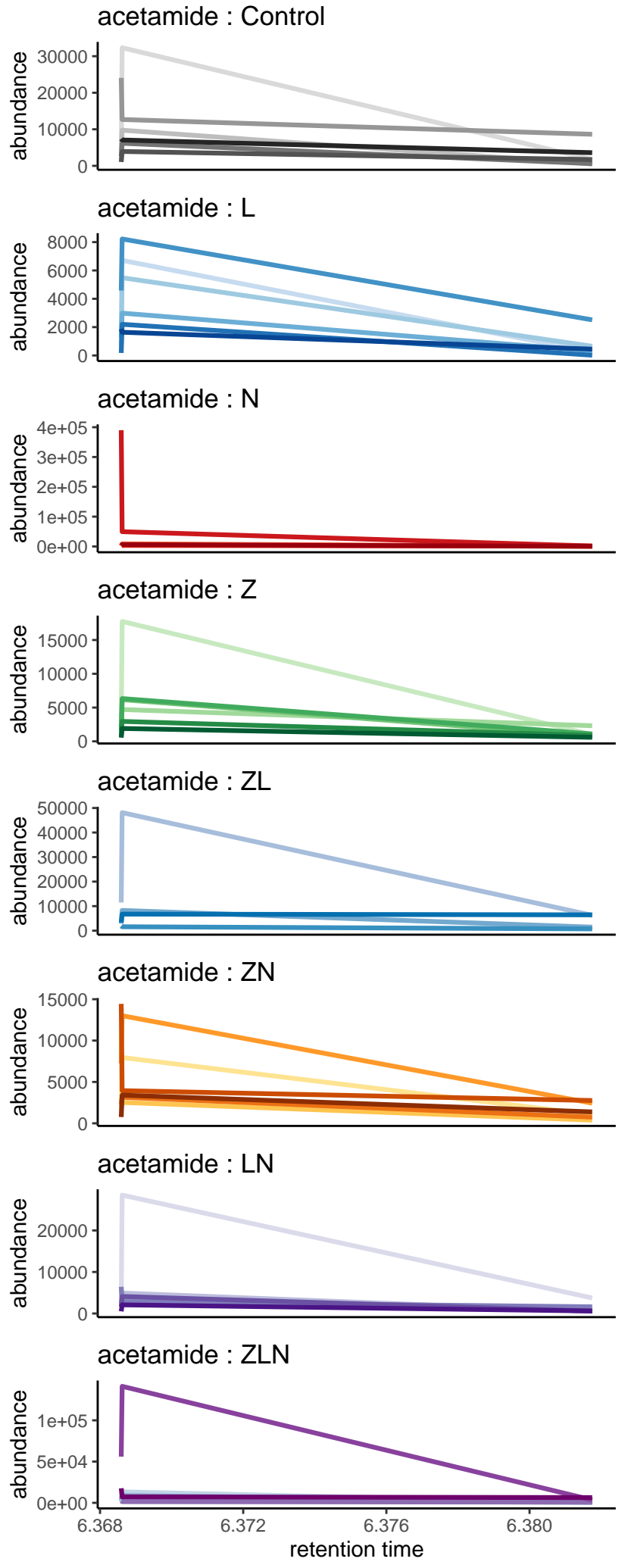


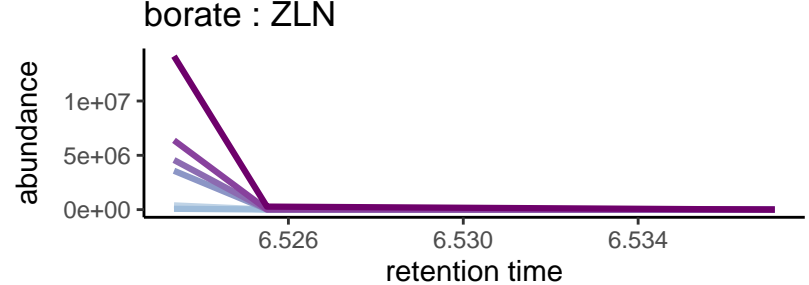
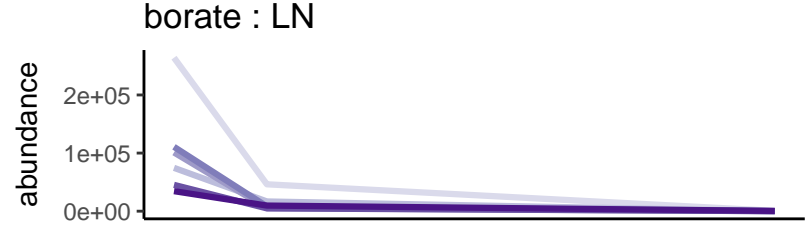
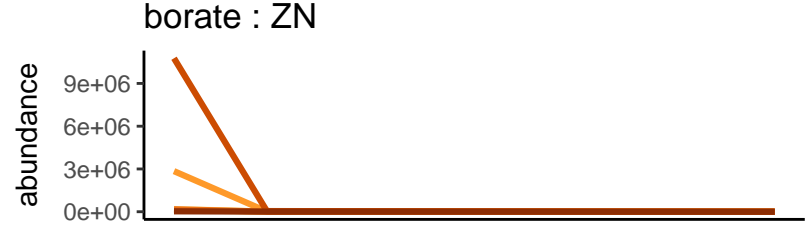
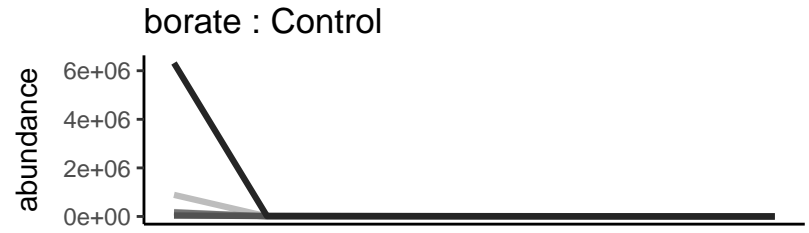
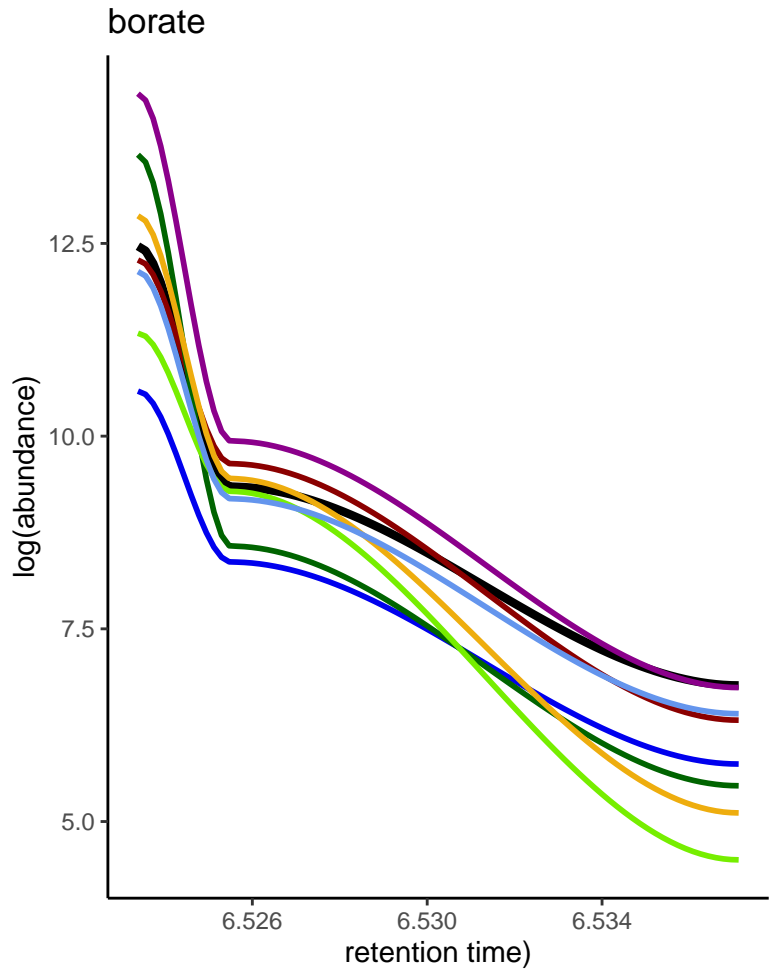


colour

c_mean	ln_mean	z_mean	zln_mean
l_mean	n_mean	zl_mean	zn_mean

acetamide								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
104	6.368592	0.61	0.61	0.26	0.03	0.79	0.95	0.54
108	6.368617	0.93	0.70	0.82	0.03	0.92	0.65	0.91
120	6.381750	0.07	0.78	0.80	0.11	0.80	0.23	0.02



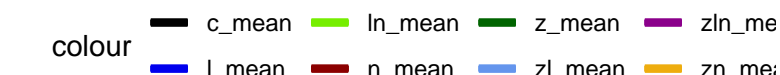


colour

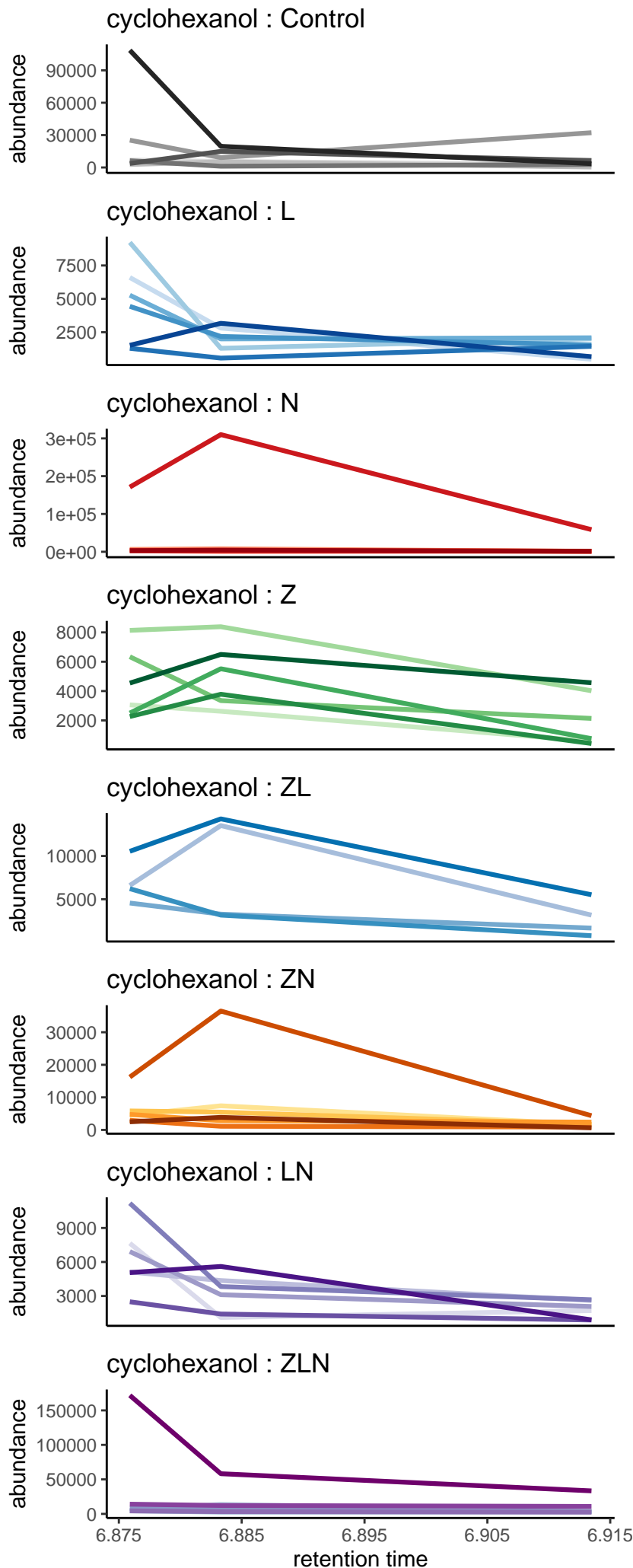
c\_mean ln\_mean z\_mean zln\_me

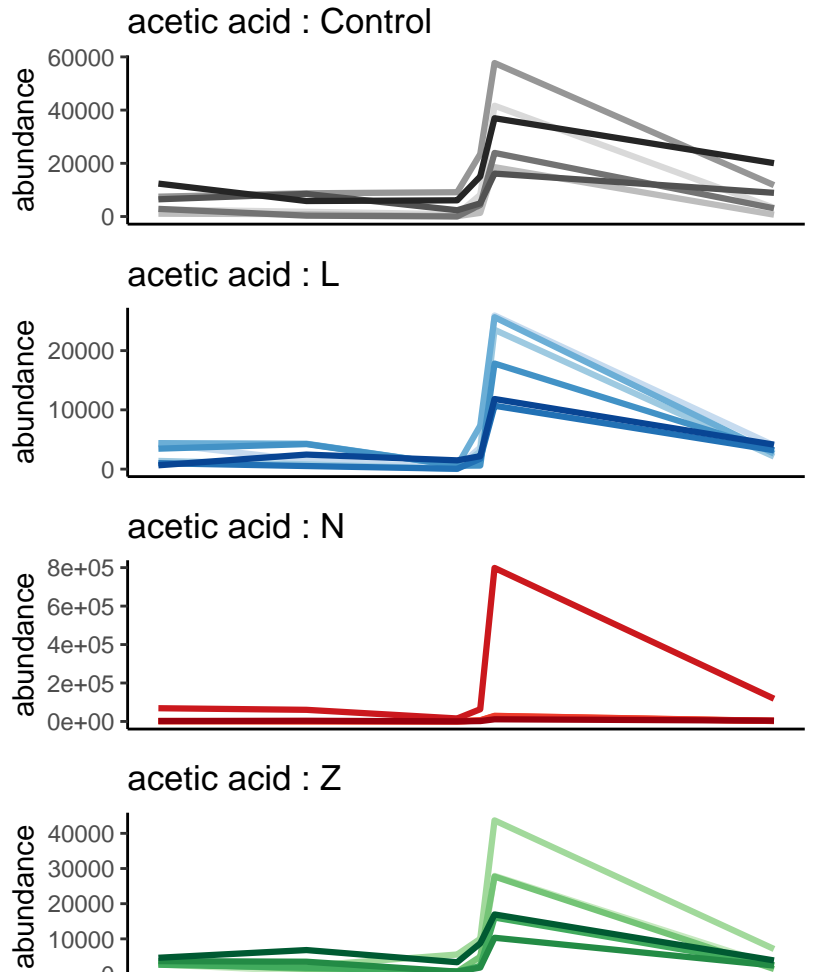
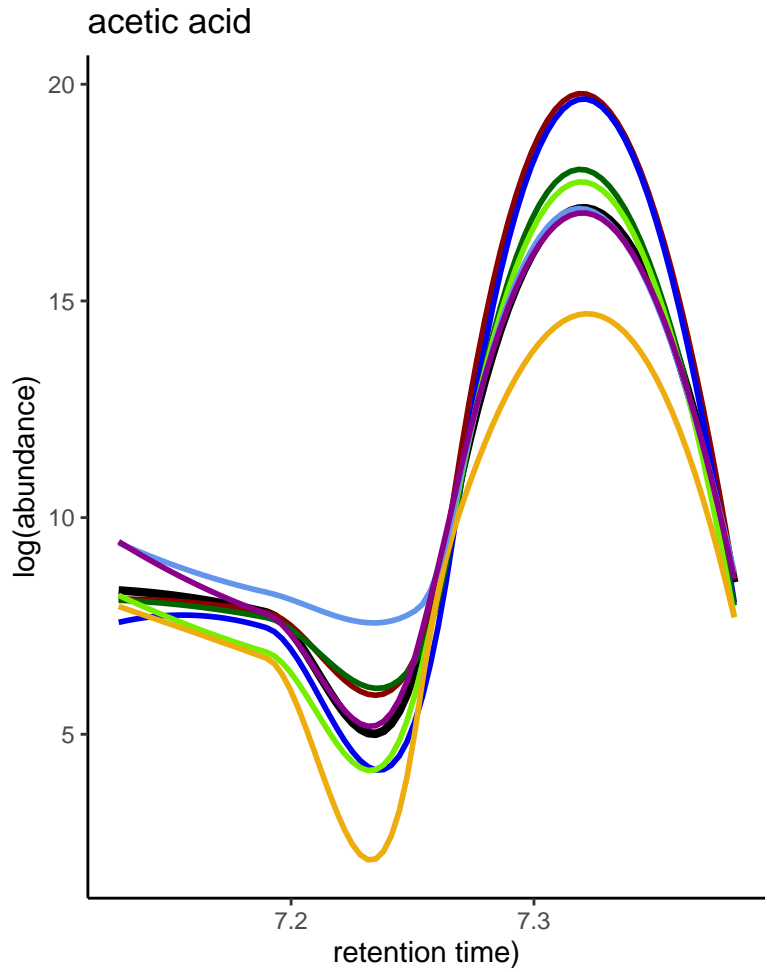
l\_mean n\_mean zl\_mean zn\_me

borate								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
125	6.523383	0.00	0.28	0.40	0.15	0.79	0.09	0.34
126	6.525508	0.66	0.87	0.02	0.06	0.74	0.67	0.56
128	6.537125	0.91	0.82	0.42	0.01	0.42	0.92	0.46



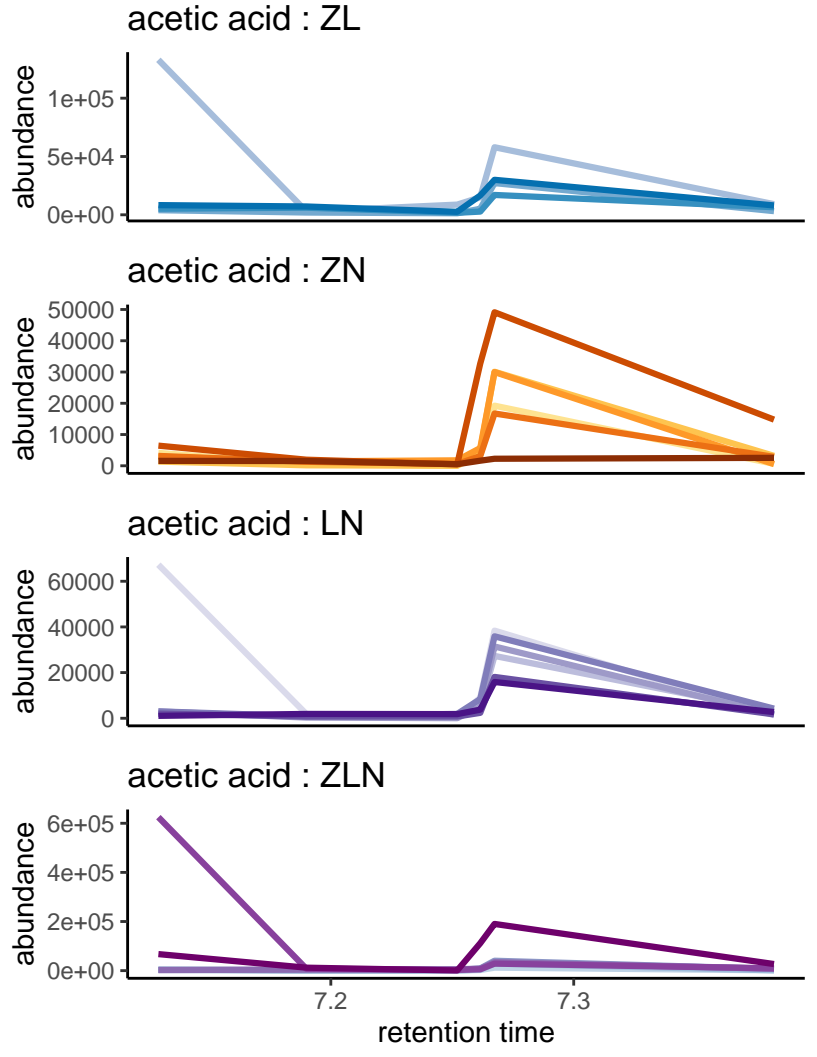
cyclohexanol								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
139	6.875892	0.91	0.97	0.49	0.04	0.63	0.37	0.68
140	6.883300	0.18	0.24	0.38	0.01	0.88	0.67	0.91
141	6.913475	0.71	0.85	0.40	0.01	0.39	0.23	0.93

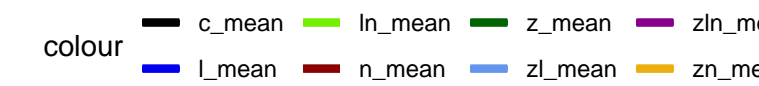
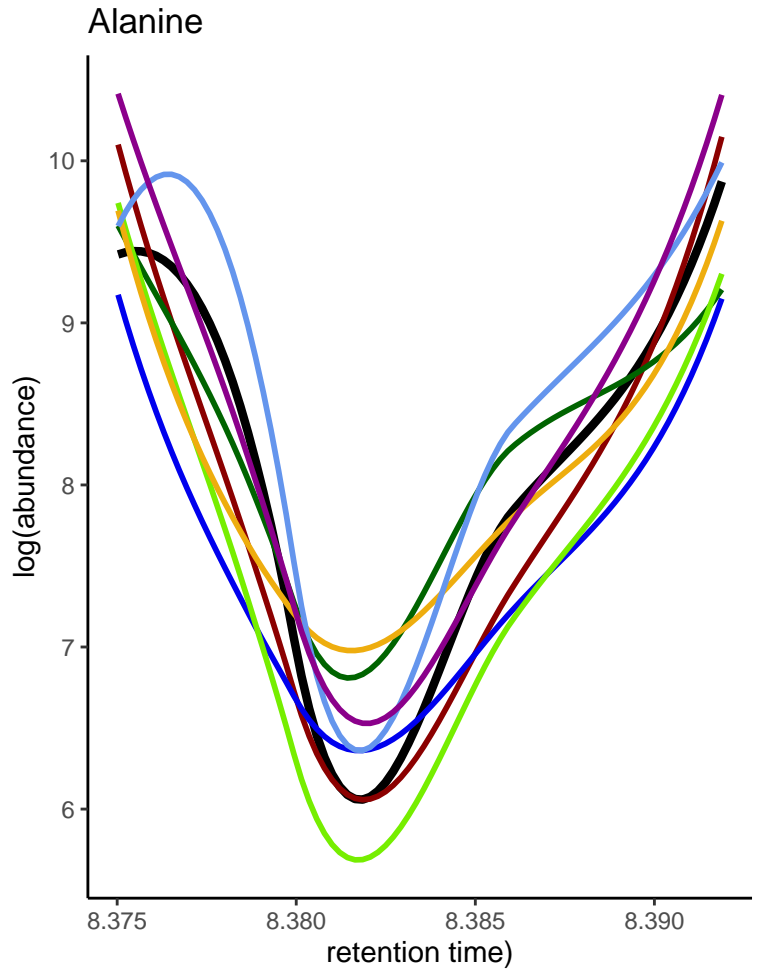




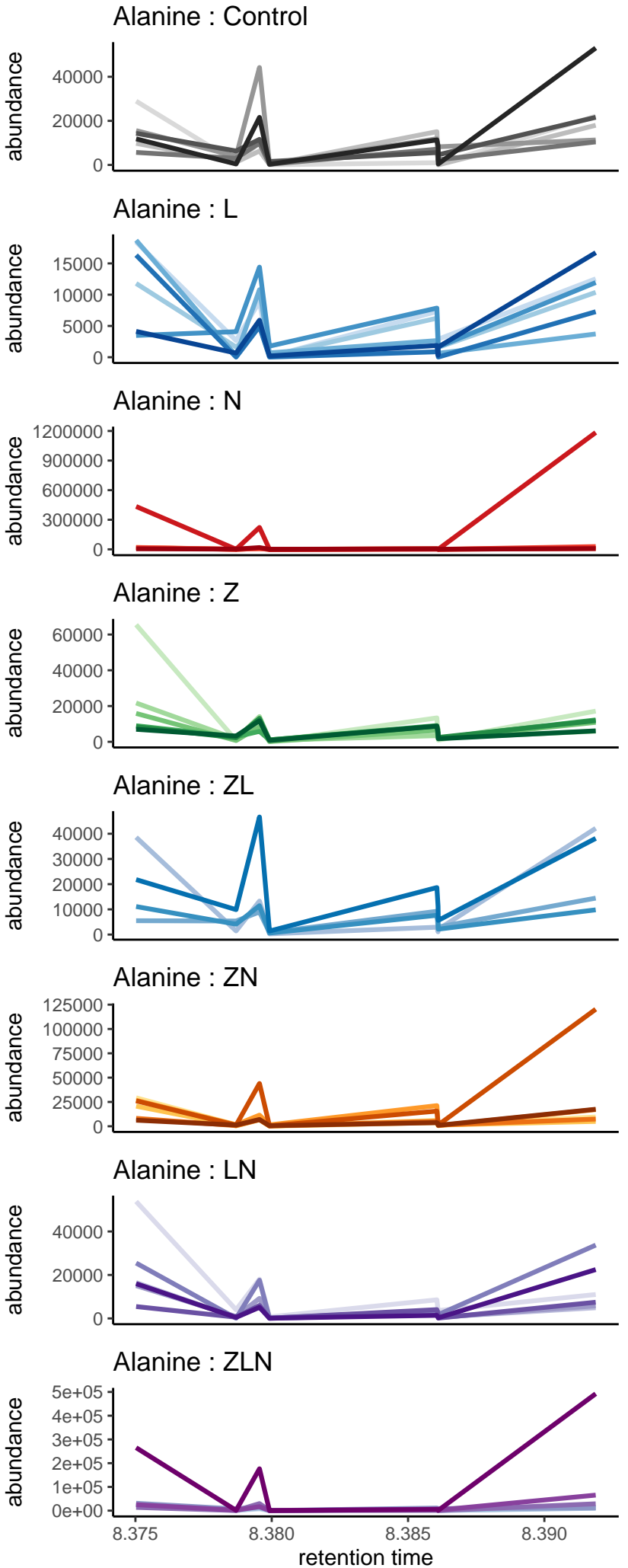
colour    c\_mean    ln\_mean    z\_mean    zln\_mean  
 l\_mean    n\_mean    zl\_mean    zn\_mean

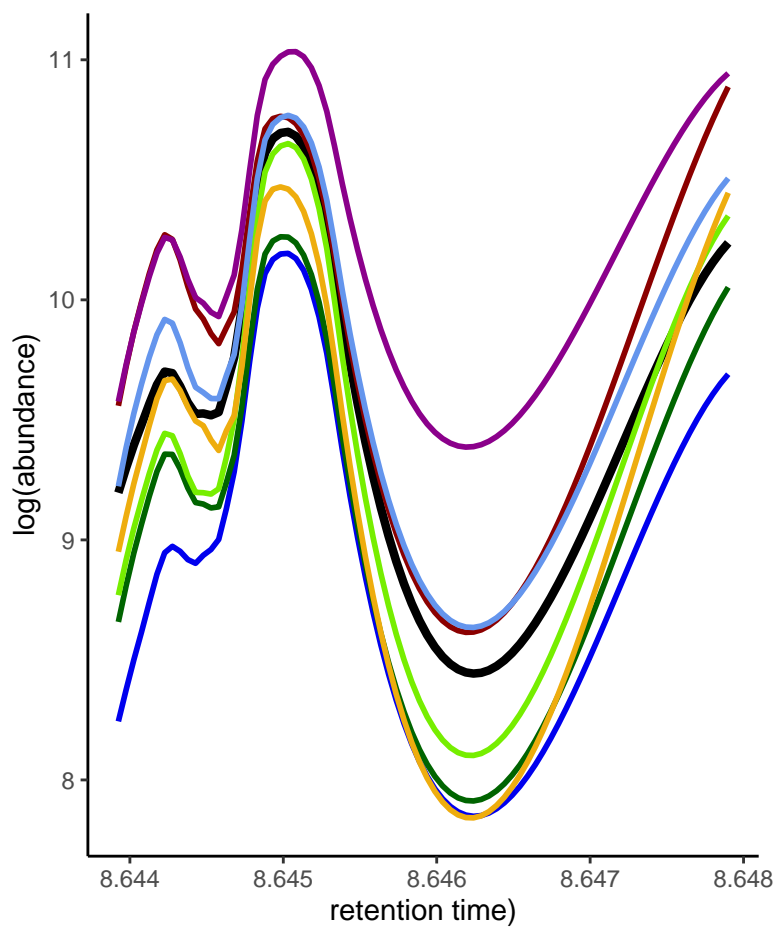
acetic acid								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
142	7.129017	0.13	0.20	0.75	0.03	0.72	0.51	0.68
155	7.189967	0.83	0.94	0.17	0.04	0.49	0.89	0.46
159	7.252033	0.45	0.86	0.62	0.03	0.12	0.65	0.97
163	7.261517	0.41	0.88	0.30	0.04	0.76	0.51	0.48
165	7.267400	0.70	0.93	0.38	0.05	0.46	0.68	0.71
166	7.382450	0.98	0.89	0.65	0.02	0.88	0.80	0.66





Alanine								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zn_fvalue
191	8.375025	0.39	0.85	0.05	0.20	0.70	0.60	0.44
192	8.378692	0.05	0.73	0.07	0.05	0.71	0.83	0.27
193	8.379550	0.45	0.95	0.39	0.02	0.93	0.71	0.81
194	8.379917	0.01	0.44	0.05	0.57	0.71	0.70	0.54
197	8.386058	0.04	0.09	0.27	0.52	0.97	0.92	0.41
198	8.386100	0.01	0.24	0.06	0.25	0.57	0.17	0.54
199	8.386108	0.05	0.96	0.28	0.51	0.50	0.79	0.81
200	8.391883	0.59	0.95	0.28	0.02	0.75	0.92	0.93

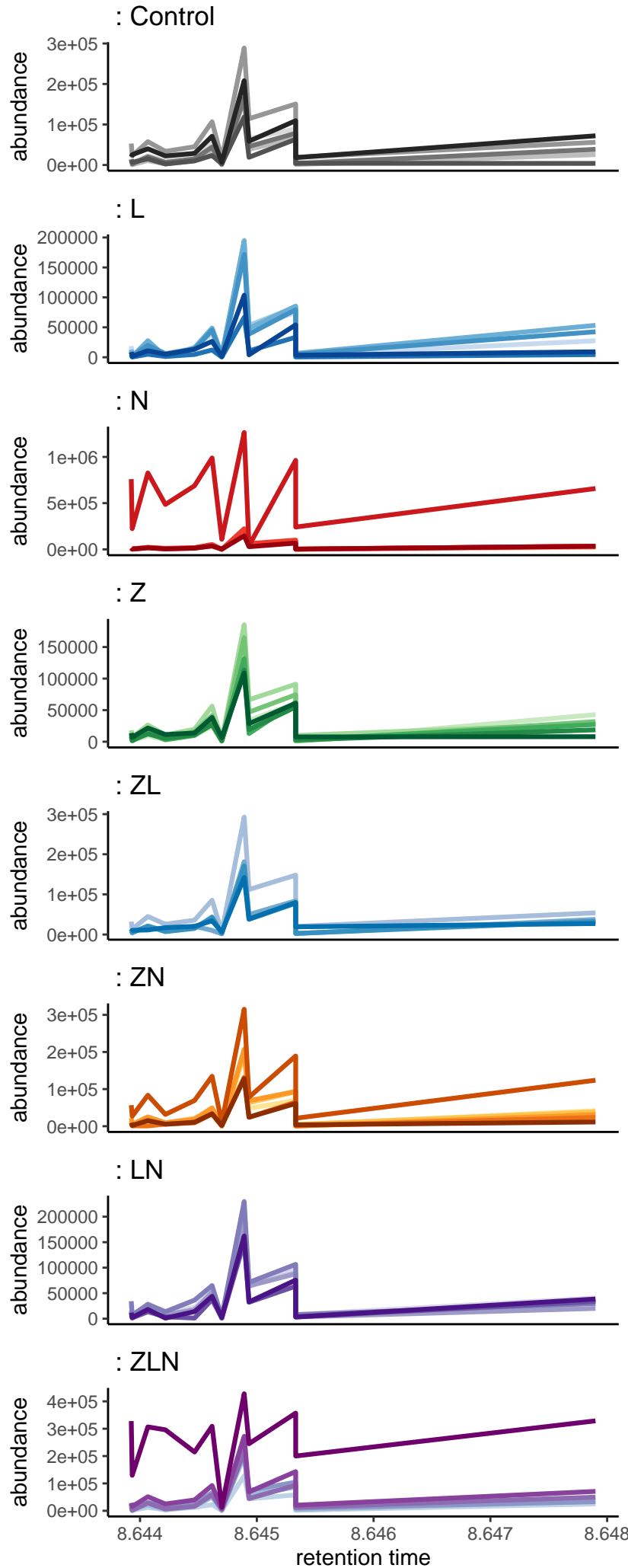




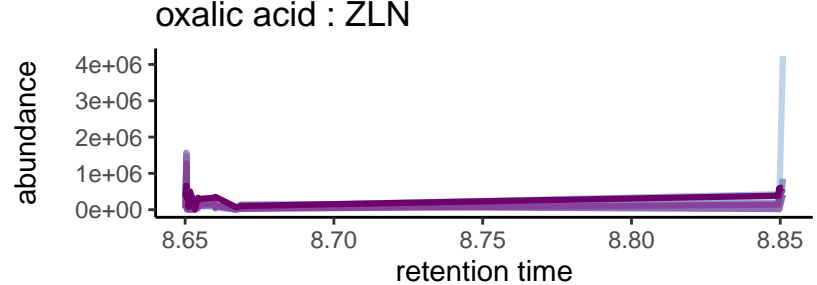
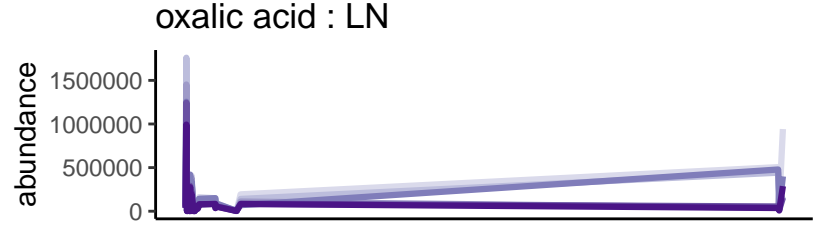
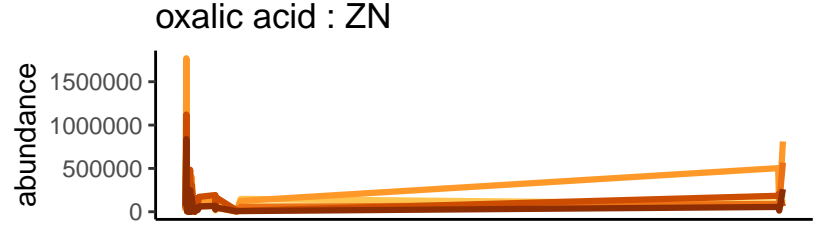
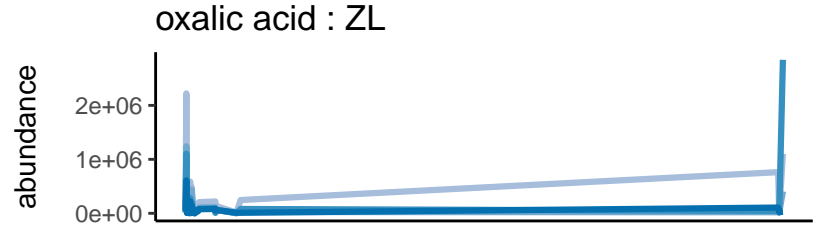
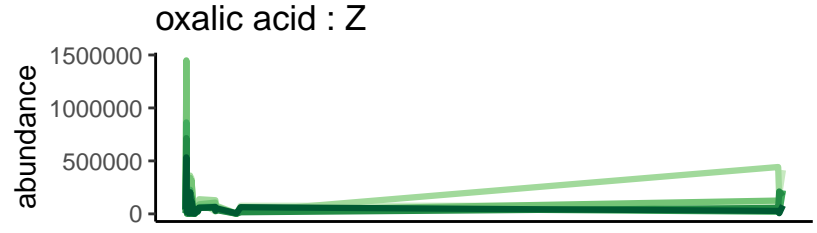
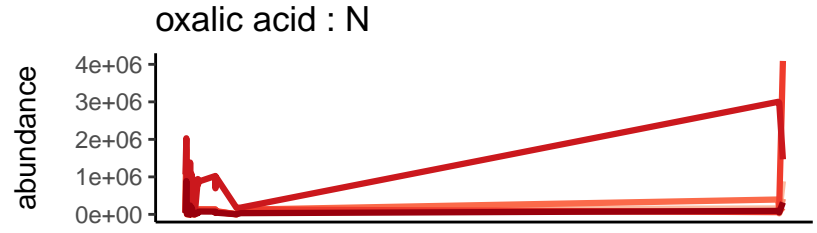
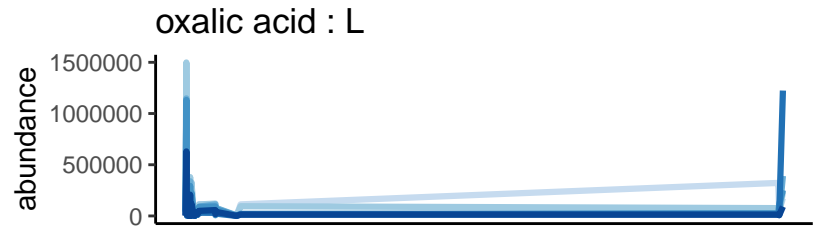
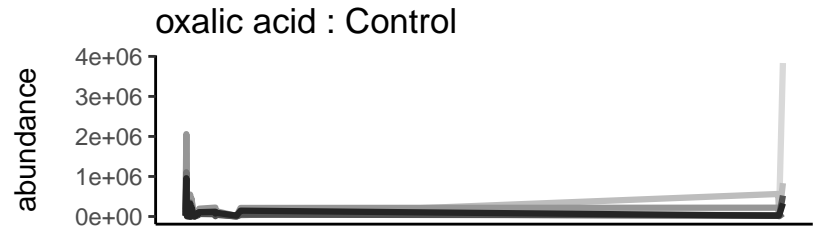
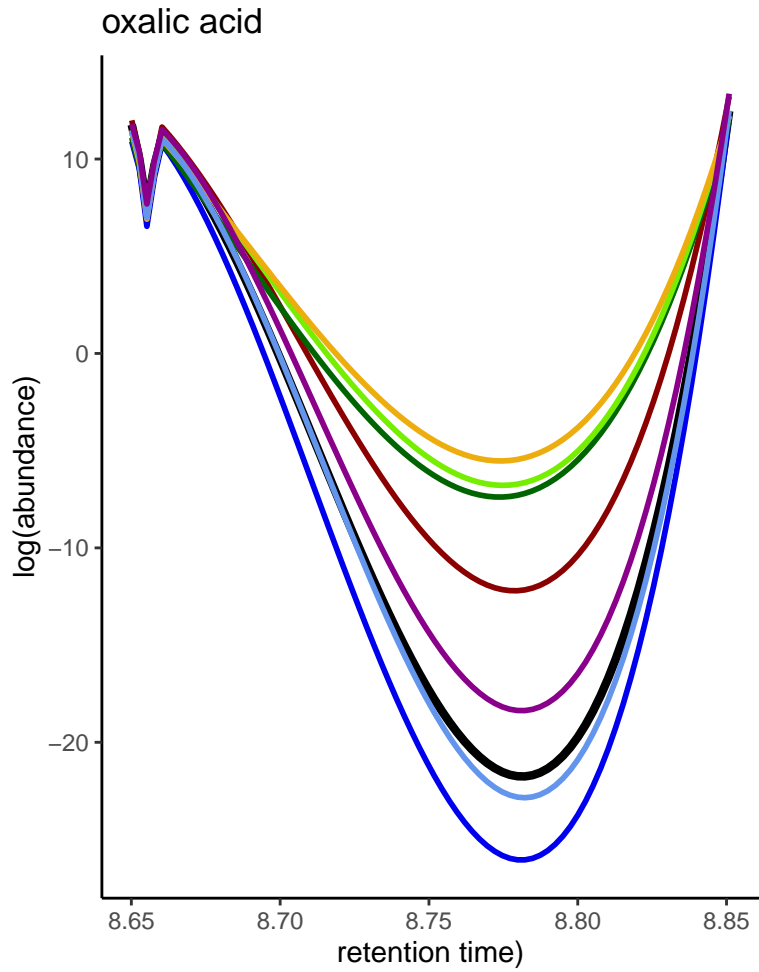
colour

c\_mean ln\_mean z\_mean zln\_me  
l\_mean n\_mean zl\_mean zn\_me

X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
207	8.643925	0.84	0.92	0.04	0.12	0.89	0.78	0.56
208	8.643933	0.34	0.24	0.44	0.01	0.68	0.92	0.44
209	8.644067	0.78	0.81	0.25	0.05	0.72	0.60	0.53
211	8.644217	0.25	0.50	0.09	0.02	0.54	0.95	0.70
213	8.644467	0.55	0.77	0.26	0.04	0.79	0.66	0.74
215	8.644617	0.75	0.69	0.03	0.17	0.85	0.60	0.64
216	8.644700	0.58	0.74	0.94	0.05	0.86	0.82	0.84
217	8.644892	0.73	0.97	0.03	0.02	0.75	0.80	0.97
219	8.644933	0.48	0.64	0.04	0.03	0.99	0.36	0.23
220	8.645333	0.99	0.92	0.05	0.04	0.85	0.85	1.00
221	8.645333	0.69	0.32	0.57	0.04	0.92	0.36	0.97
223	8.647900	0.45	0.89	0.03	0.06	0.64	0.96	0.98





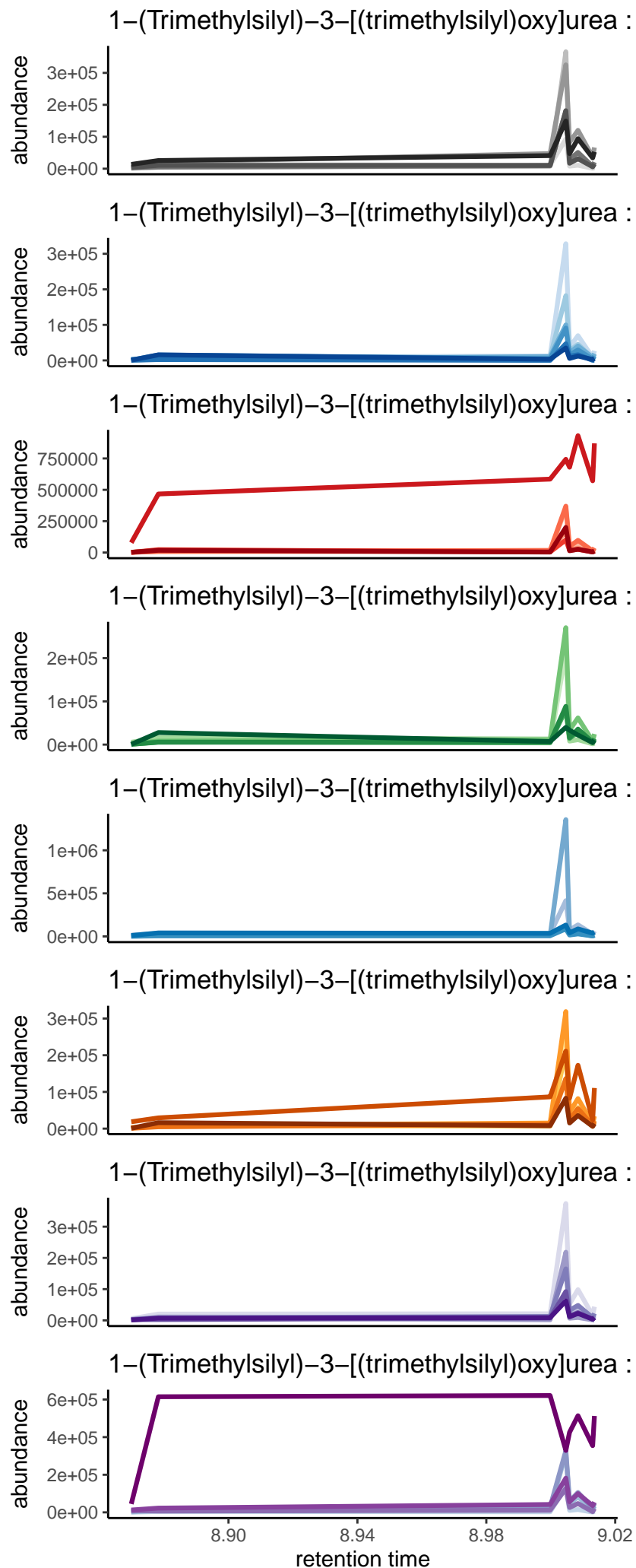


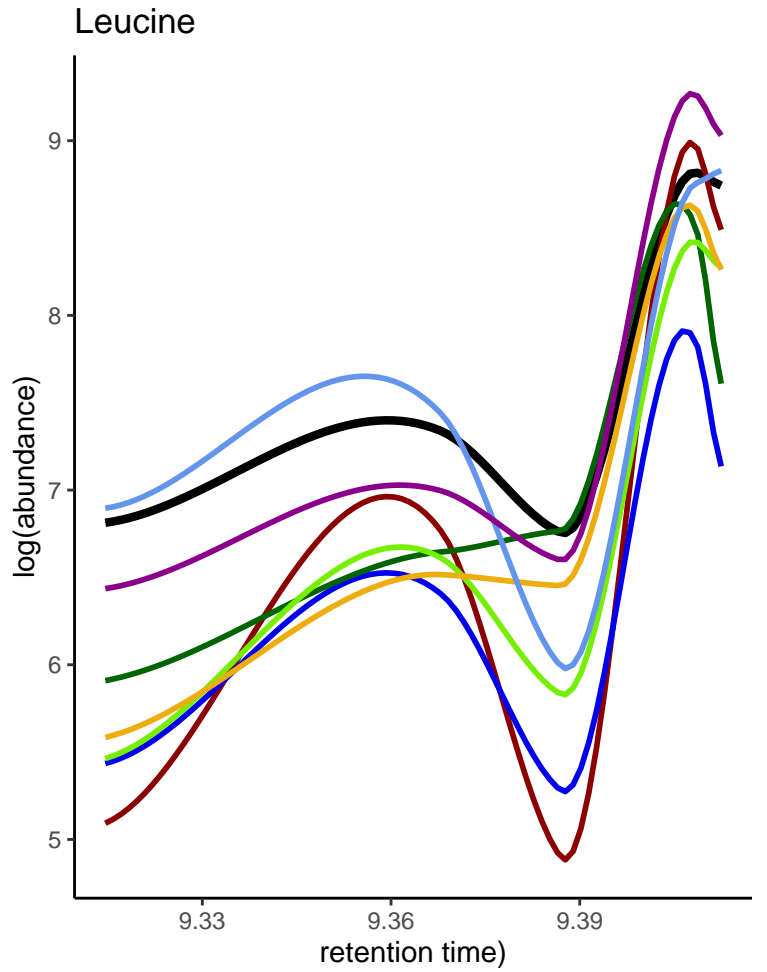
colour

c\_mean ln\_mean z\_mean zln\_mean  
l\_mean n\_mean zl\_mean zn\_mean

oxalic acid										
X	retention_time	z_value	l_value	n_value	zl_value	zn_value	ln_value	zn_value	zl_value	zn_value
231	8.659133	0.64	0.41	0.07	0.05	0.72	0.43	0.57		
234	8.650417	0.29	0.96	0.03	0.16	0.50	0.78	0.45		
235	8.650642	0.32	0.84	0.68	0.03	0.30	0.53	0.63		
236	8.650658	0.94	0.83	0.50	0.01	0.69	0.86	0.15		
238	8.651175	0.78	0.93	0.03	0.04	0.66	0.98	0.90		
237	8.651175	0.76	0.84	0.05	0.03	0.90	0.64	0.72		
239	8.651175	0.76	0.93	0.15	0.05	0.95	0.79	0.91		
242	8.651392	0.52	0.64	0.16	0.04	0.39	0.67	0.58		
243	8.651417	0.87	0.94	0.03	0.02	0.76	0.91	0.98		
244	8.651483	0.59	0.70	0.06	0.02	0.78	0.84	0.81		
245	8.651483	0.29	0.93	0.19	0.03	0.89	0.74	0.80		
248	8.651592	0.86	0.99	0.11	0.05	0.86	0.88	0.88		
249	8.651617	0.90	0.91	0.03	0.03	0.68	0.99	0.92		
250	8.651617	0.84	0.96	0.04	0.05	0.79	0.98	1.00		
257	8.651667	0.99	0.74	0.04	0.06	0.96	0.70	0.81		
259	8.651667	0.75	0.91	0.02	0.02	0.94	0.75	0.67		
256	8.651667	0.33	0.93	0.23	0.01	0.44	0.97	0.61		
258	8.651667	0.82	0.98	0.06	0.04	0.81	0.96	0.92		
261	8.651683	0.53	0.96	0.03	0.05	0.60	0.96	0.80		
262	8.651692	0.60	0.56	0.09	0.02	0.97	0.66	0.84		
265	8.651757	0.65	0.70	0.31	0.04	0.94	0.78	0.56		
266	8.651767	0.70	0.89	0.24	0.02	0.71	0.58	0.88		
267	8.651783	0.85	0.98	0.03	0.03	0.76	0.93	0.94		
268	8.651783	0.82	0.89	0.05	0.04	0.67	0.89	0.90		
280	8.651783	0.70	0.62	0.05	0.31	0.85	0.87	0.36		
273	8.651783	0.87	0.99	0.08	0.05	0.79	0.75	0.99		
282	8.651808	0.74	0.90	0.03	0.03	0.61	0.92	0.88		
283	8.651808	0.89	0.58	0.10	0.04	0.73	0.68	0.92		
293	8.651900	0.70	0.83	0.23	0.04	0.94	0.89	0.96		
294	8.651917	0.71	0.74	0.15	0.04	0.68	0.70	0.85		
299	8.651925	0.57	0.87	0.24	0.04	0.92	0.71	0.68		
300	8.651933	0.70	0.87	0.26	0.02	0.60	0.53	0.80		
305	8.651967	0.92	0.81	0.09	0.03	0.70	0.90	0.83		
310	8.652008	0.92	0.87	0.03	0.04	0.65	0.83	0.87		
318	8.652150	0.92	0.84	0.03	0.04	0.76	0.93	0.96		
319	8.652150	0.77	0.81	0.05	0.03	0.72	1.00	0.91		
320	8.652150	0.65	0.69	0.03	0.06	0.51	0.93	0.49		
321	8.652790	0.84	0.94	0.03	0.05	0.64	1.00	0.85		
322	8.652790	0.68	0.98	0.35	0.03	0.89	0.60	0.96		
324	8.653250	0.74	0.85	0.34	0.37	0.60	0.01	0.71		
326	8.654283	0.94	0.90	0.04	0.03	0.82	0.89	0.99		
327	8.654283	1.00	0.94	0.06	0.03	0.76	1.00	0.93		
329	8.654700	0.77	0.77	0.02	0.03	0.58	0.90	0.87		
333	8.660183	0.50	0.71	0.02	0.02	0.75	0.76	0.77		
334	8.660192	0.39	0.70	0.03	0.17	0.48	0.26	0.31		
335	8.660383	0.99	0.93	0.05	0.05	0.79	0.91	0.95		
340	8.662783	0.55	0.93	0.13	0.03	0.89	0.95	0.96		
341	8.668842	0.13	0.83	0.04	0.08	0.38	0.15	0.30		
349	8.849300	0.56	0.82	0.01	0.43	0.34	0.96	0.94		
350	8.849650	0.65	0.25	0.03	0.52	0.70	0.43	0.16		
351	8.850883	0.86	0.51	0.19	0.01	0.81	0.78	0.42		

1-(Trimethylsilyl)-3-[(trimethylsilyl)oxy]urea								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
358	8.869833	0.55	0.76	0.17	0.00	0.92	0.49	0.27
359	8.878242	0.50	0.61	0.21	0.04	0.57	0.80	0.66
360	8.999800	0.26	0.65	0.18	0.02	0.83	0.64	0.48
362	9.004683	0.73	0.79	0.11	0.01	0.68	0.45	0.22
363	9.005833	0.30	0.84	0.14	0.03	0.72	0.56	0.99
364	9.008450	0.37	0.78	0.09	0.03	0.48	0.37	0.93
366	9.013000	0.38	0.72	0.14	0.03	0.94	0.71	0.65
367	9.013550	0.20	0.49	0.12	0.03	0.74	0.64	0.84

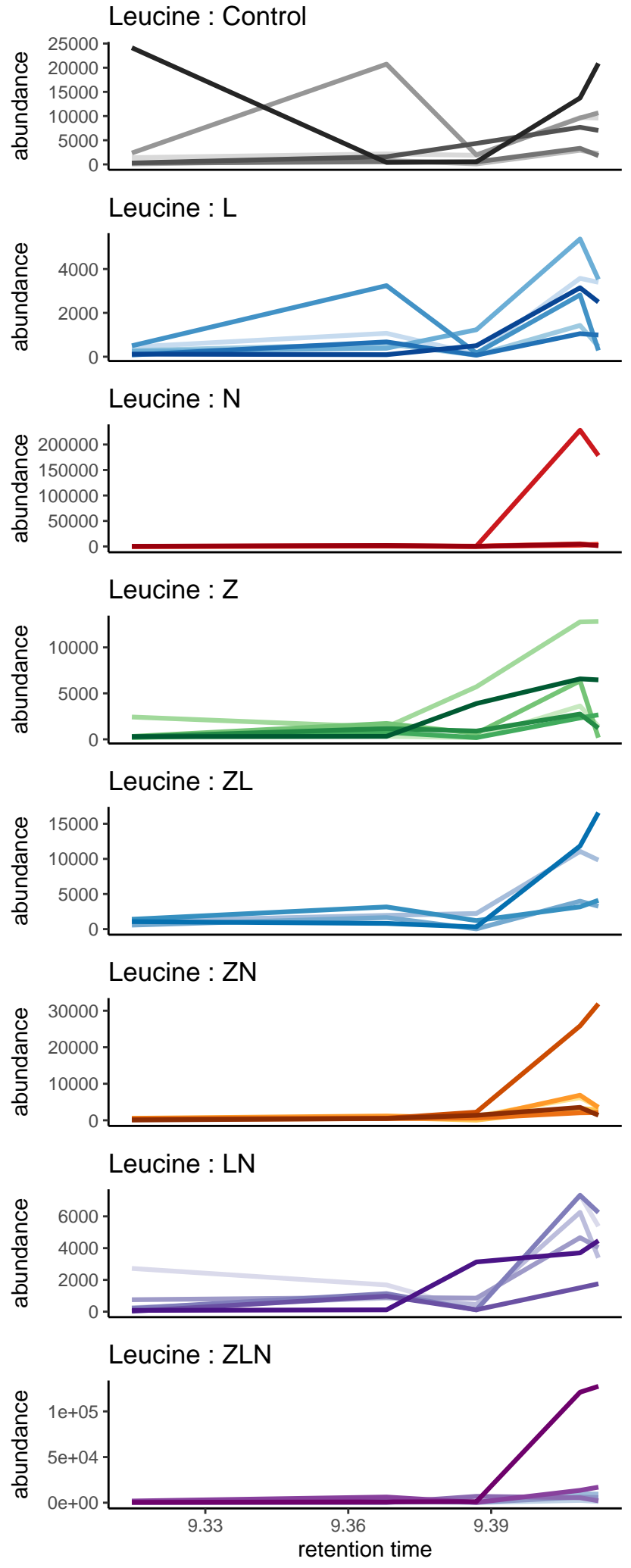


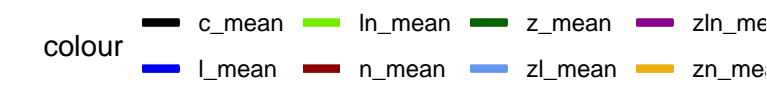
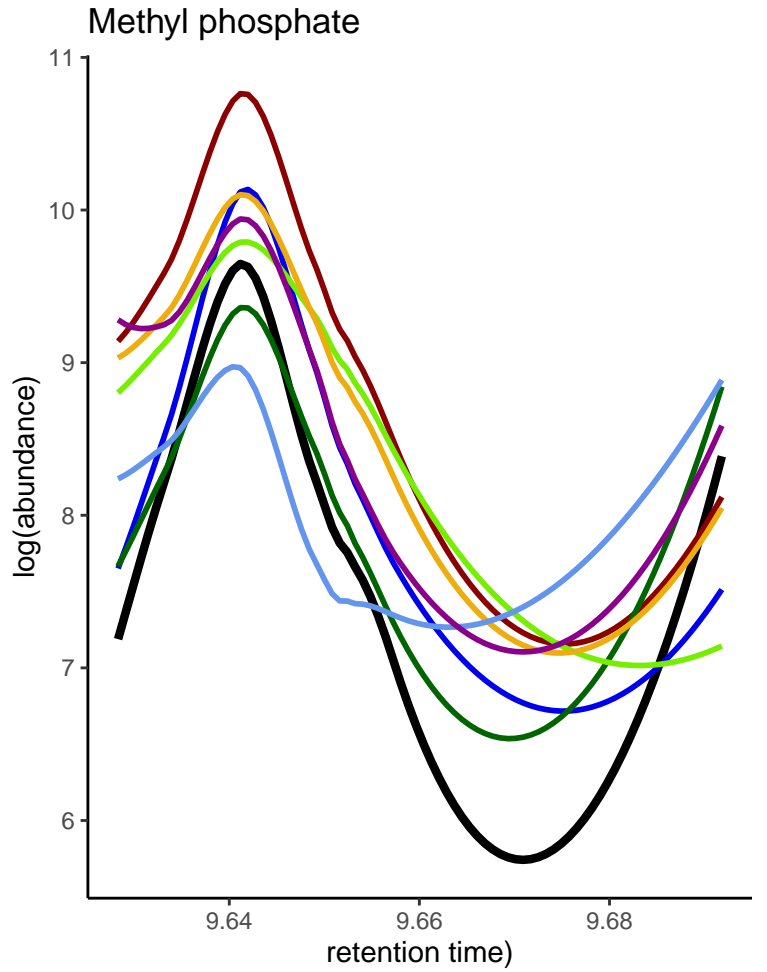


colour

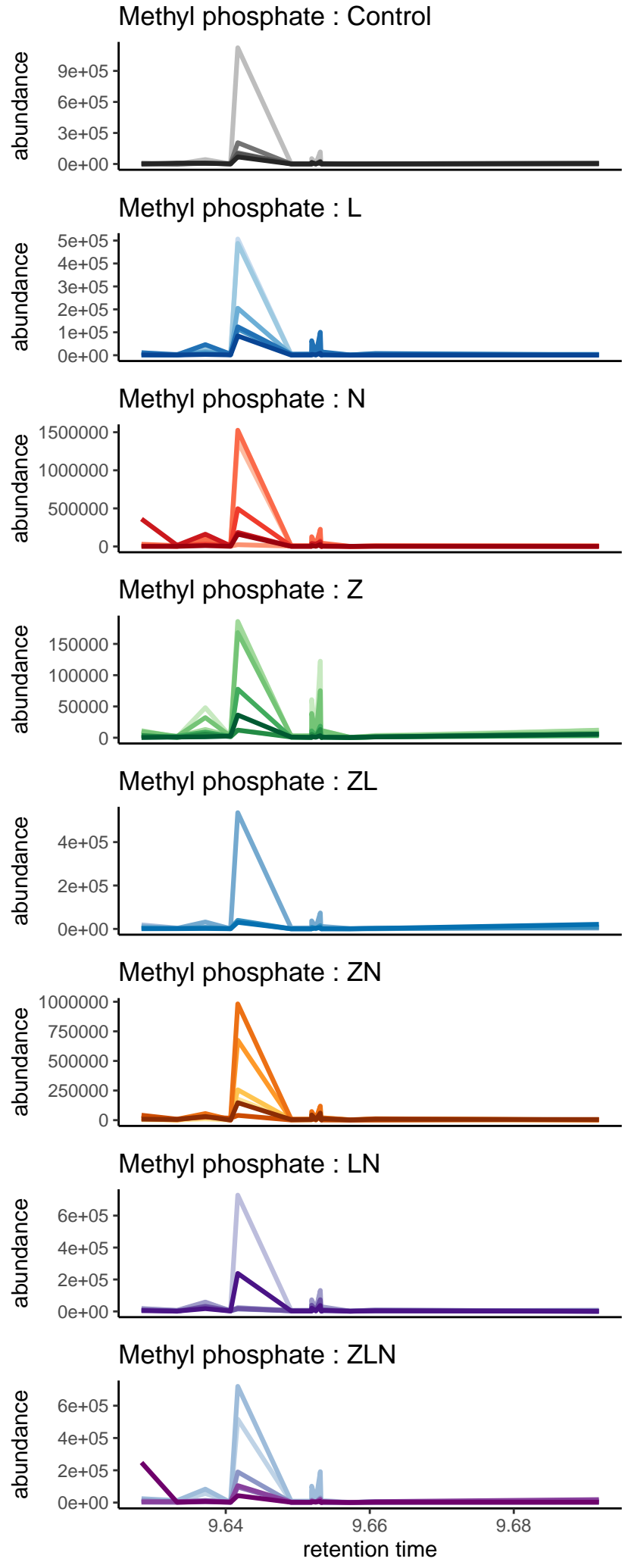
c_mean	ln_mean	z_mean	zln_mean
l_mean	n_mean	zl_mean	zn_mean

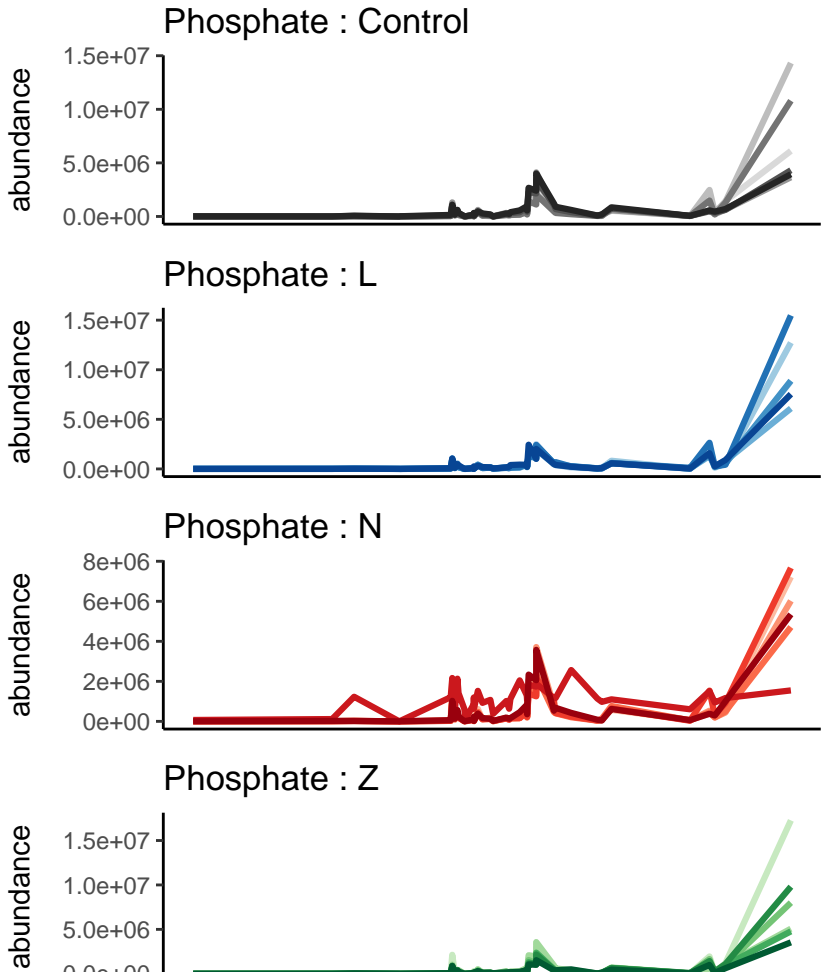
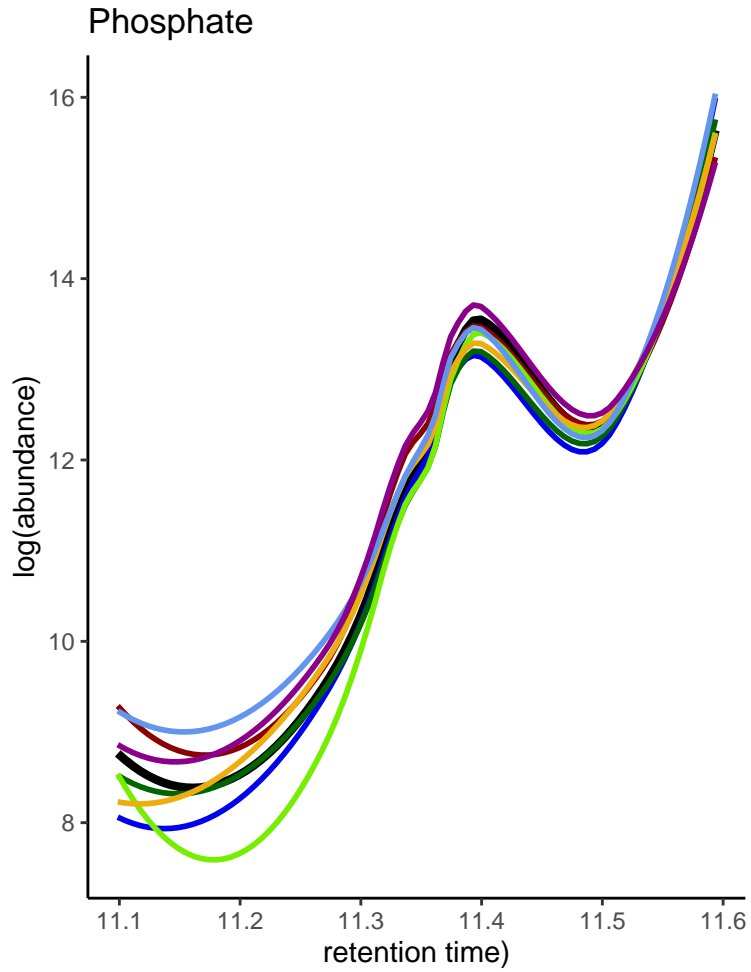
Leucine								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
371	9.314583	0.21	0.67	0.10	0.05	0.51	0.23	0.19
377	9.368025	0.80	0.99	0.49	0.05	0.88	0.64	0.34
378	9.386892	0.06	0.53	0.48	0.95	0.37	0.05	0.38
385	9.408650	0.35	0.60	0.22	0.04	0.90	0.48	0.95
391	9.412500	0.53	0.99	0.19	0.02	0.99	0.49	0.22





Methyl phosphate								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zn_fvalue
394	9.628300	0.44	0.57	0.00	0.70	0.70	0.41	0.61
395	9.633242	0.39	0.51	0.00	0.63	0.99	0.64	0.77
396	9.637167	0.11	0.38	0.00	0.43	0.83	0.27	0.52
397	9.640667	0.05	0.88	0.02	0.32	0.24	0.93	0.02
398	9.641667	0.44	0.38	0.22	0.27	0.03	0.09	0.40
399	9.649083	0.22	0.74	0.00	0.08	0.91	0.73	0.11
400	9.651933	0.18	1.00	0.00	0.09	0.32	0.58	0.14
401	9.651933	0.18	0.72	0.00	0.31	0.79	0.32	0.65
402	9.652517	0.31	0.87	0.00	0.91	0.22	0.32	0.25
403	9.653133	0.15	0.59	0.01	0.35	0.90	0.33	0.79
404	9.653200	0.09	0.26	0.00	0.14	0.65	0.86	0.61
405	9.653308	0.38	0.70	0.00	0.03	0.45	0.12	0.70
406	9.653433	0.29	0.39	0.00	0.13	0.72	0.39	0.54
407	9.657283	0.37	0.22	0.00	0.58	0.25	0.20	0.81
408	9.660767	0.86	0.25	0.00	0.17	0.32	0.10	0.56
410	9.691817	0.00	0.13	0.09	0.01	0.59	0.71	0.53

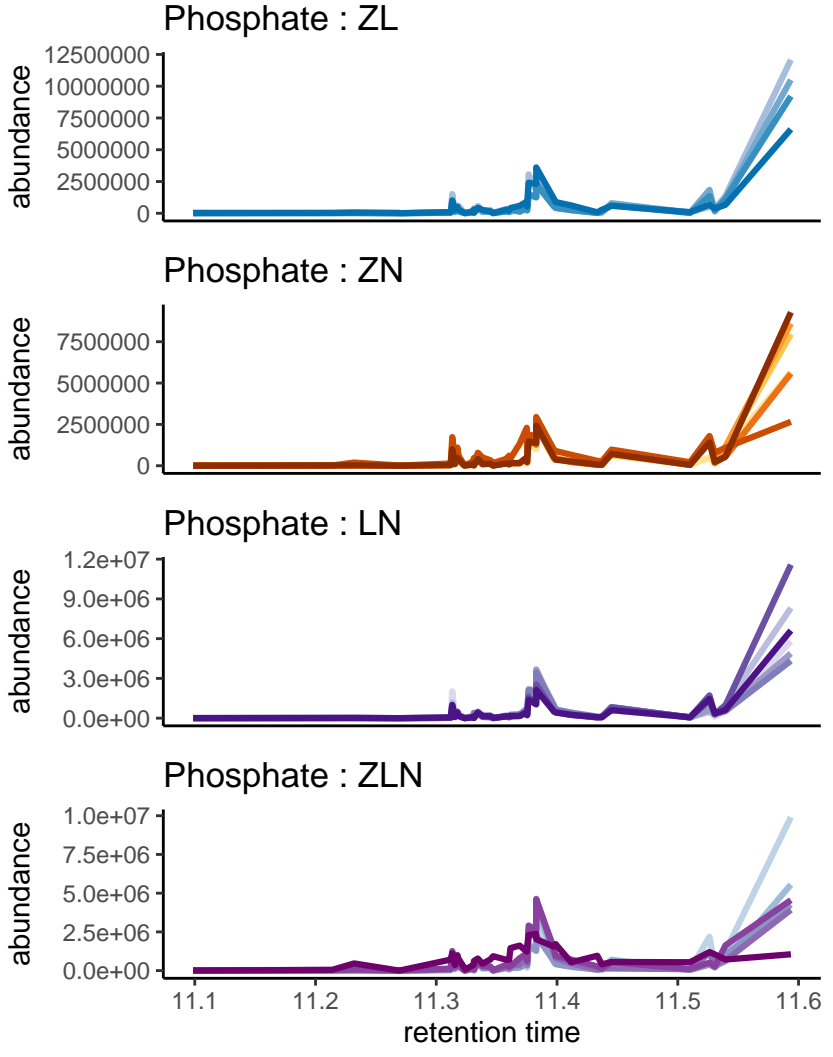


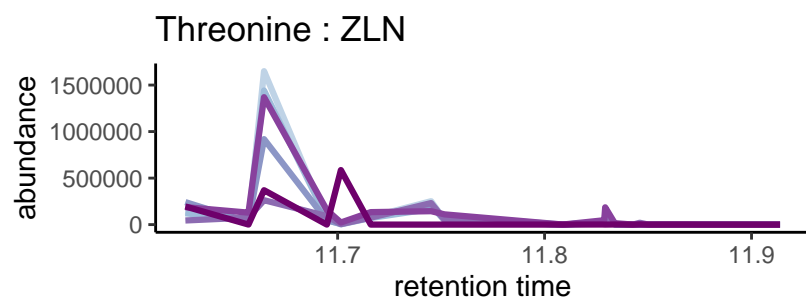
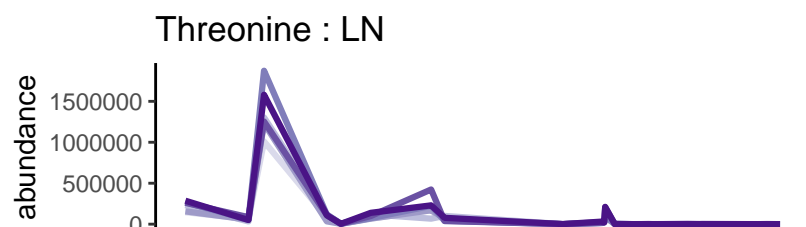
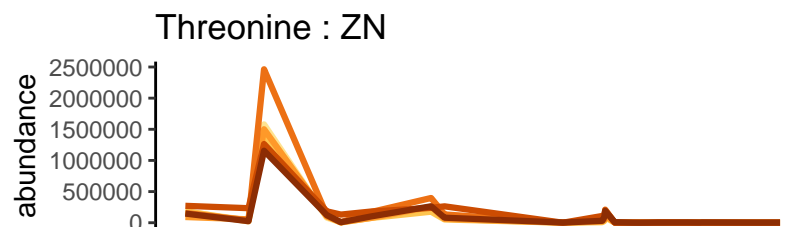
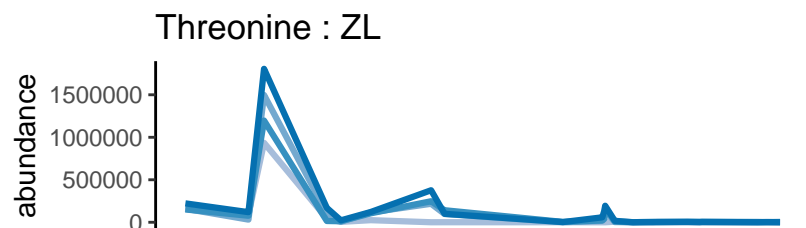
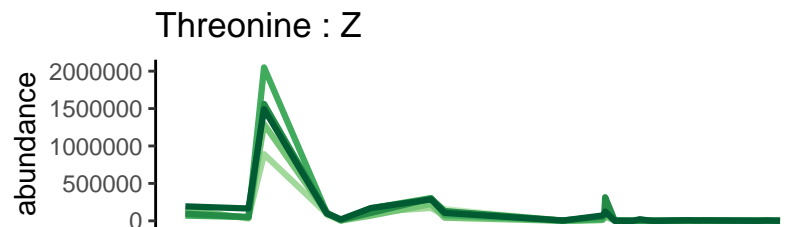
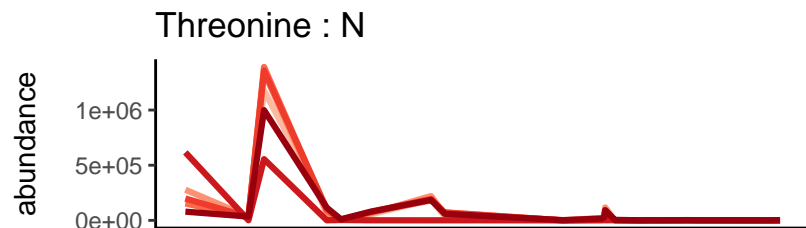
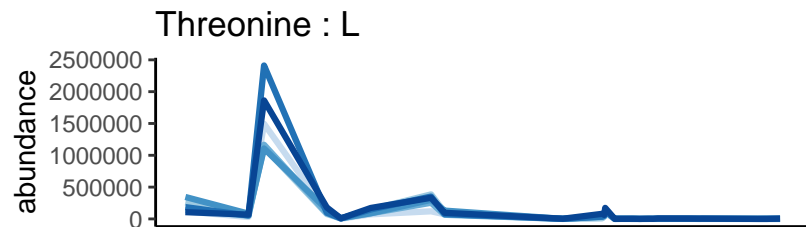
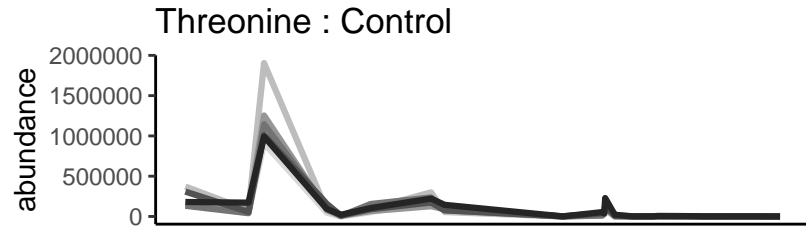
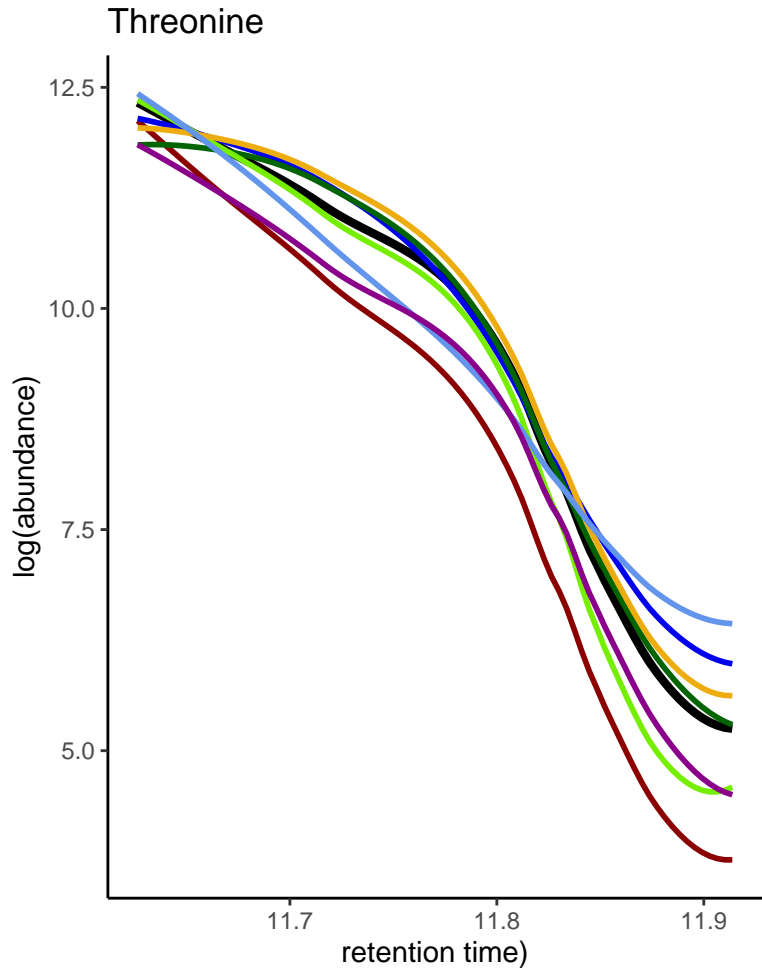


colour

- c\_mean
- ln\_mean
- z\_mean
- zln\_mean
- l\_mean
- n\_mean
- zl\_mean
- zn\_mean

Phosphate									
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue	
417	11.09869	0.97	0.80	0.77	0.04	0.20	0.87	0.96	
421	11.21377	0.16	0.92	0.08	0.04	0.62	0.71	0.99	
425	11.23197	0.13	0.15	0.63	0.04	0.30	0.21	0.19	
448	11.26938	0.03	0.45	0.16	0.84	0.33	0.04	0.99	
449	11.27008	0.69	0.27	0.69	0.02	0.60	0.71	0.97	
497	11.31195	0.43	0.92	0.97	0.05	0.45	0.72	0.34	
499	11.31325	0.89	0.21	0.03	0.59	0.79	0.95	0.69	
504	11.31538	0.57	0.89	0.45	0.04	0.73	0.53	0.67	
507	11.31752	0.92	0.92	0.03	0.02	0.25	0.18	0.60	
509	11.31842	0.07	0.80	0.05	0.51	0.21	0.51	0.83	
514	11.32328	0.50	0.90	0.11	0.03	0.75	0.44	0.97	
515	11.32398	0.24	0.21	0.04	0.04	0.73	0.87	0.96	
519	11.33110	0.95	0.63	0.09	0.04	0.60	0.95	0.98	
520	11.33110	0.89	1.00	0.13	0.03	0.81	0.87	0.98	
521	11.33117	0.41	0.55	0.24	0.04	0.91	0.59	0.94	
523	11.33445	0.78	0.63	0.01	0.01	0.37	0.86	0.82	
524	11.33823	0.50	0.20	0.10	0.02	0.77	0.84	0.84	
527	11.34479	0.36	0.17	0.05	0.26	0.83	0.49	0.11	
532	11.34707	0.47	0.62	0.98	1.00	0.44	0.02	0.05	
540	11.35803	0.48	0.80	0.31	0.00	0.62	0.49	0.94	
543	11.36035	0.76	0.84	0.35	0.04	0.93	0.97	1.00	
546	11.36141	0.59	0.52	0.80	0.02	0.75	0.61	0.75	
551	11.36910	0.82	0.74	0.44	0.03	0.75	0.83	0.56	
553	11.37493	0.62	0.89	0.02	0.04	0.65	0.70	0.72	
554	11.37520	0.50	0.60	0.26	0.01	0.61	0.39	0.83	
555	11.37648	0.68	0.19	0.41	0.00	0.53	0.28	0.30	
560	11.38278	0.87	0.64	0.11	0.00	0.85	0.05	0.41	
561	11.38278	0.90	0.65	0.24	0.04	0.81	0.07	0.21	
564	11.39743	0.66	0.38	0.08	0.01	0.54	0.21	0.68	
565	11.39883	0.88	0.27	0.28	0.02	0.59	0.47	0.90	
571	11.41173	0.85	0.56	0.27	0.02	0.37	0.60	0.72	
576	11.43338	0.94	0.85	0.01	0.30	0.12	0.49	0.47	
577	11.43708	0.97	0.93	1.00	0.04	0.85	0.99	0.97	
579	11.44505	0.21	0.04	0.89	0.14	0.45	0.13	0.03	
590	11.51013	0.23	0.61	0.02	0.20	0.61	0.49	0.92	
596	11.52625	0.52	0.61	0.13	0.01	0.22	0.24	0.80	
599	11.53063	0.31	0.50	0.23	0.03	0.54	0.78	0.91	
602	11.53948	0.02	0.35	0.38	0.11	0.67	0.30	0.36	
614	11.59367	0.65	0.45	0.02	0.14	0.66	0.21	0.34	

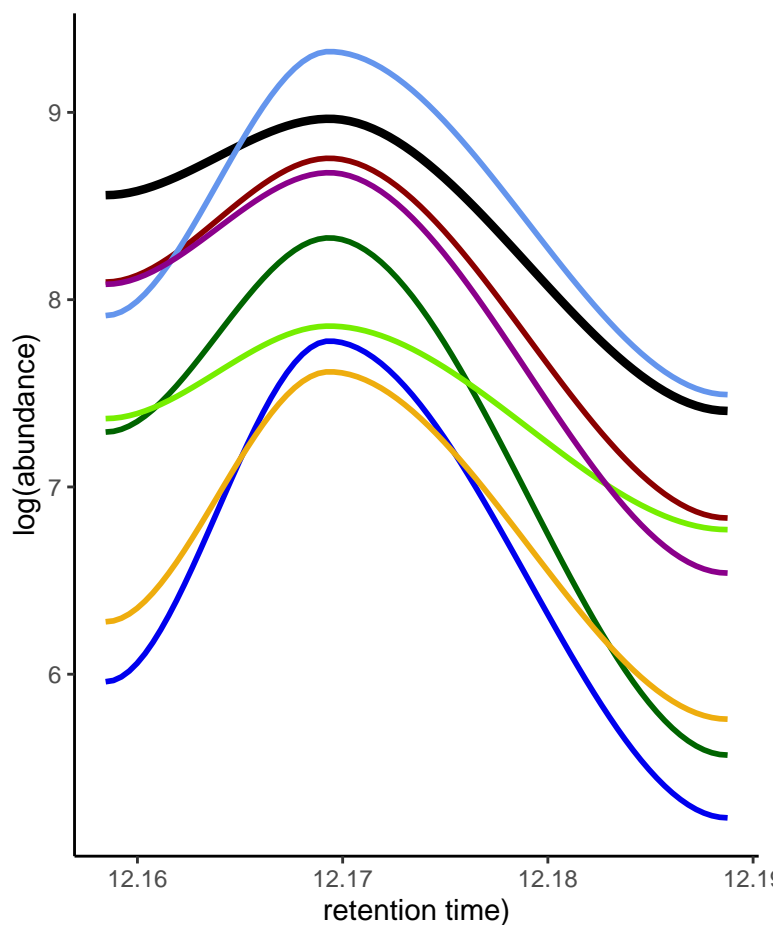




colour    c\_mean    ln\_mean    z\_mean    zln\_me  
          l\_mean    n\_mean    zl\_mean    zn\_me

Threonine								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
620	11.62642	0.02	0.85	0.44	0.70	0.87	0.54	0.19
626	11.65690	0.94	0.78	0.04	0.30	0.84	0.66	0.05
628	11.66443	0.83	0.53	0.22	0.01	0.63	0.27	0.26
647	11.69478	0.70	0.57	0.26	0.04	0.65	0.87	0.32
651	11.70157	0.03	0.43	0.96	0.89	0.31	0.74	0.71
658	11.71600	0.91	0.86	0.12	0.04	0.69	0.93	0.24
664	11.74517	0.66	0.48	0.42	0.04	0.43	0.67	0.77
666	11.75168	0.83	0.34	0.30	0.05	0.30	0.70	0.58
687	11.80883	0.14	0.58	0.03	0.47	0.53	0.44	0.43
697	11.82788	0.95	0.72	0.50	0.05	0.05	0.37	0.88
698	11.82882	0.92	0.39	0.38	0.03	0.19	0.52	0.96
699	11.82928	0.57	0.35	0.60	0.04	0.39	0.46	0.66
702	11.83392	0.05	0.52	0.70	0.24	0.90	0.65	0.04
707	11.84275	0.69	0.46	0.04	0.54	0.29	0.42	0.47
709	11.84598	0.03	0.62	0.14	0.89	0.38	0.20	0.48
711	11.85023	0.39	0.59	0.02	0.31	0.77	0.78	0.21
713	11.85240	0.18	0.69	0.01	0.17	0.15	0.18	0.26
718	11.85472	0.24	0.63	0.08	0.92	0.03	0.10	0.78
722	11.86910	0.34	0.81	0.02	0.18	0.49	0.19	0.83
727	11.87495	0.23	0.82	0.01	0.38	0.40	0.42	0.31
729	11.87605	0.12	0.56	0.00	0.19	0.28	0.32	0.56
734	11.90287	0.27	0.73	0.02	0.57	0.47	0.66	0.31
735	11.90750	0.12	0.58	0.03	0.33	0.67	0.43	0.12
736	11.91378	0.58	0.33	0.03	0.41	0.21	0.14	0.29

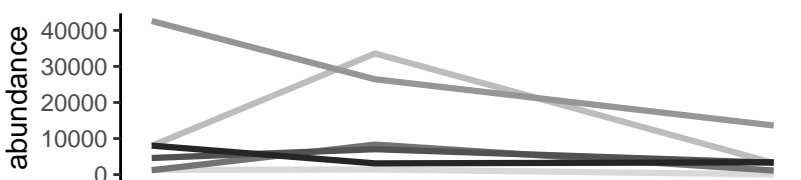
Glyceric acid



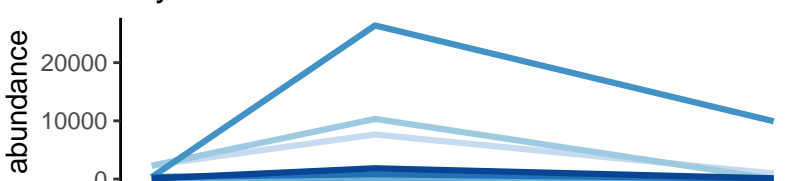
colour    c\_mean    l\_mean    n\_mean    z\_mean    zn\_mean    zl\_mean    ln\_mean    zln\_mean

Glyceric acid								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
744	12.15845	0.76	0.56	0.85	0.00	0.33	0.12	0.72
747	12.16933	0.89	0.86	0.54	0.05	0.54	0.85	0.91
750	12.18876	0.58	0.98	0.78	0.05	0.50	0.62	0.17

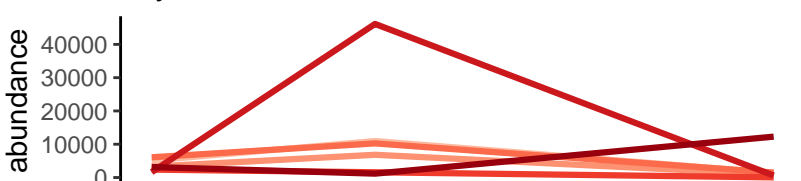
Glyceric acid : Control



Glyceric acid : L



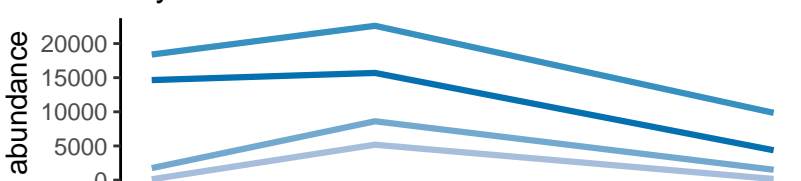
Glyceric acid : N



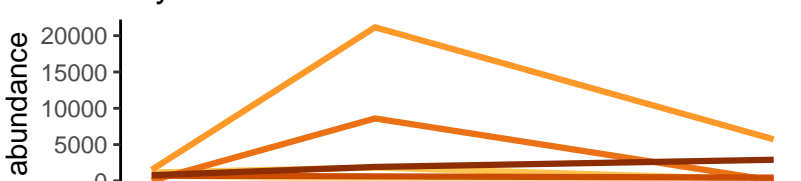
Glyceric acid : Z



Glyceric acid : ZL



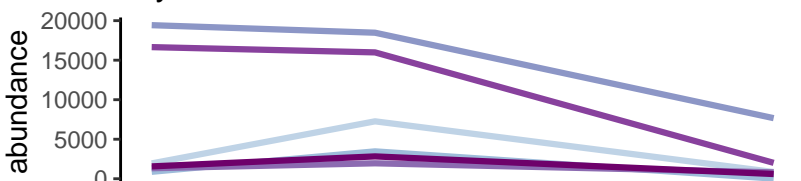
Glyceric acid : ZN



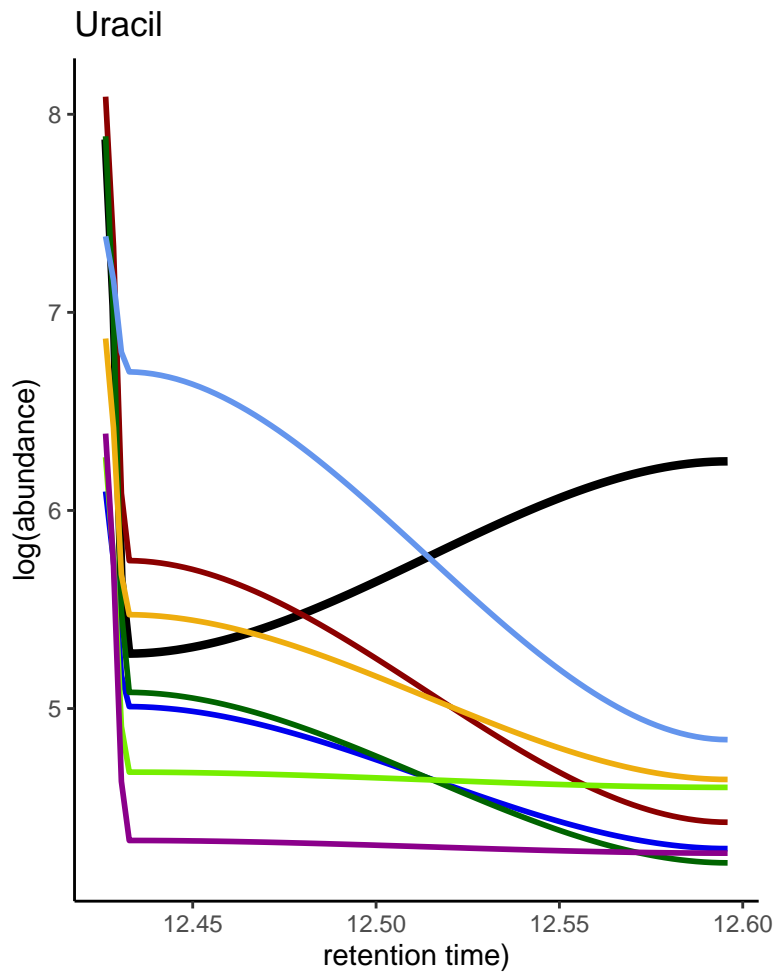
Glyceric acid : LN



Glyceric acid : ZLN



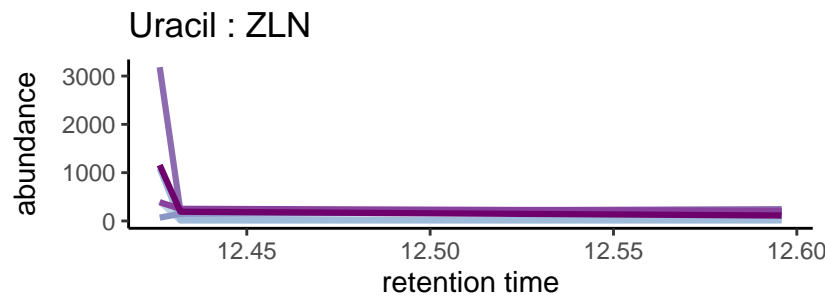
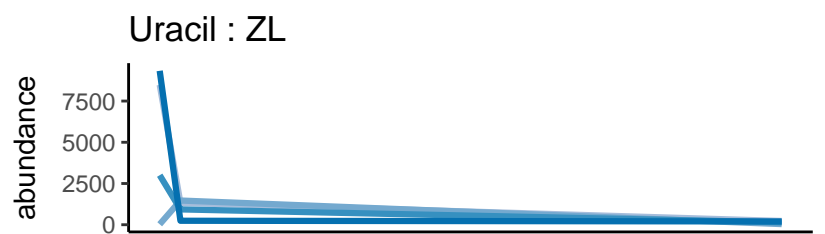
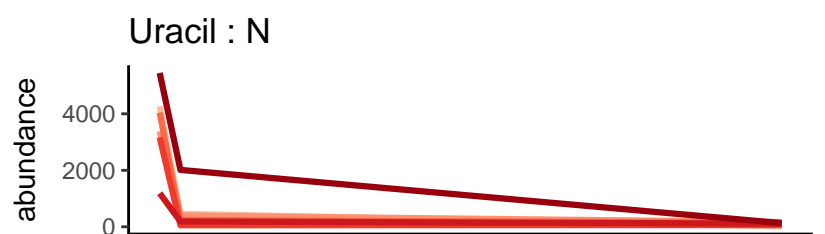
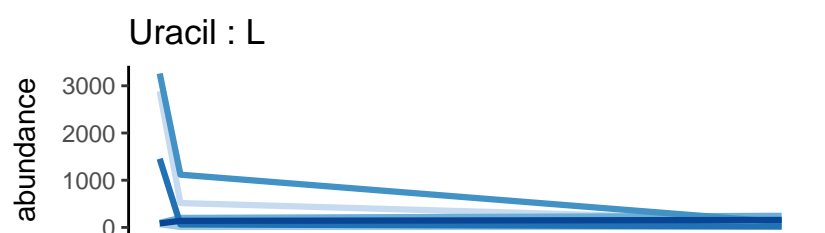
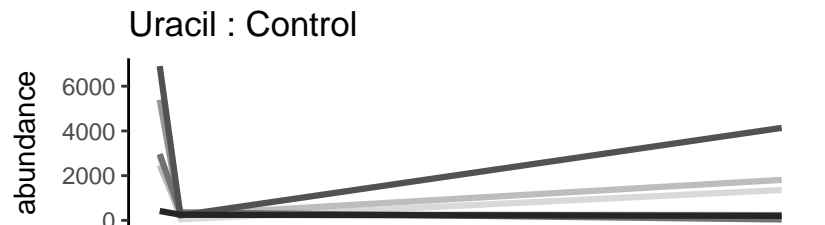
retention time



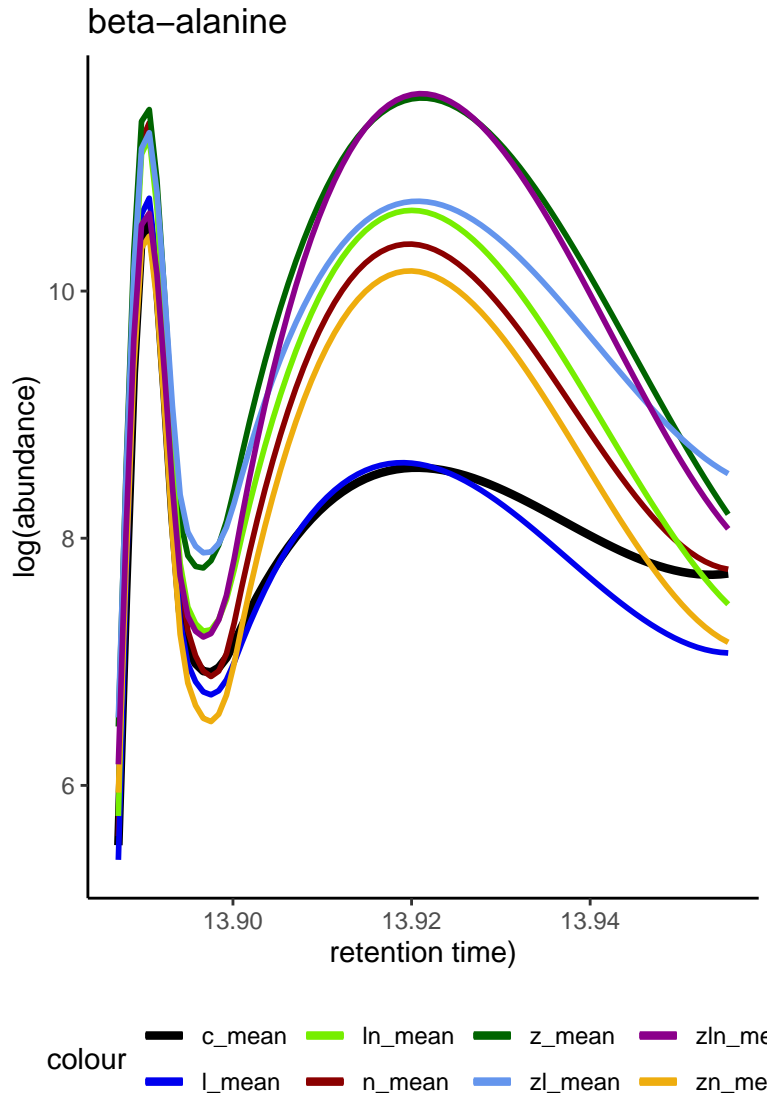
colour

c_mean	ln_mean	z_mean	zln_mean
l_mean	n_mean	zl_mean	zn_mean

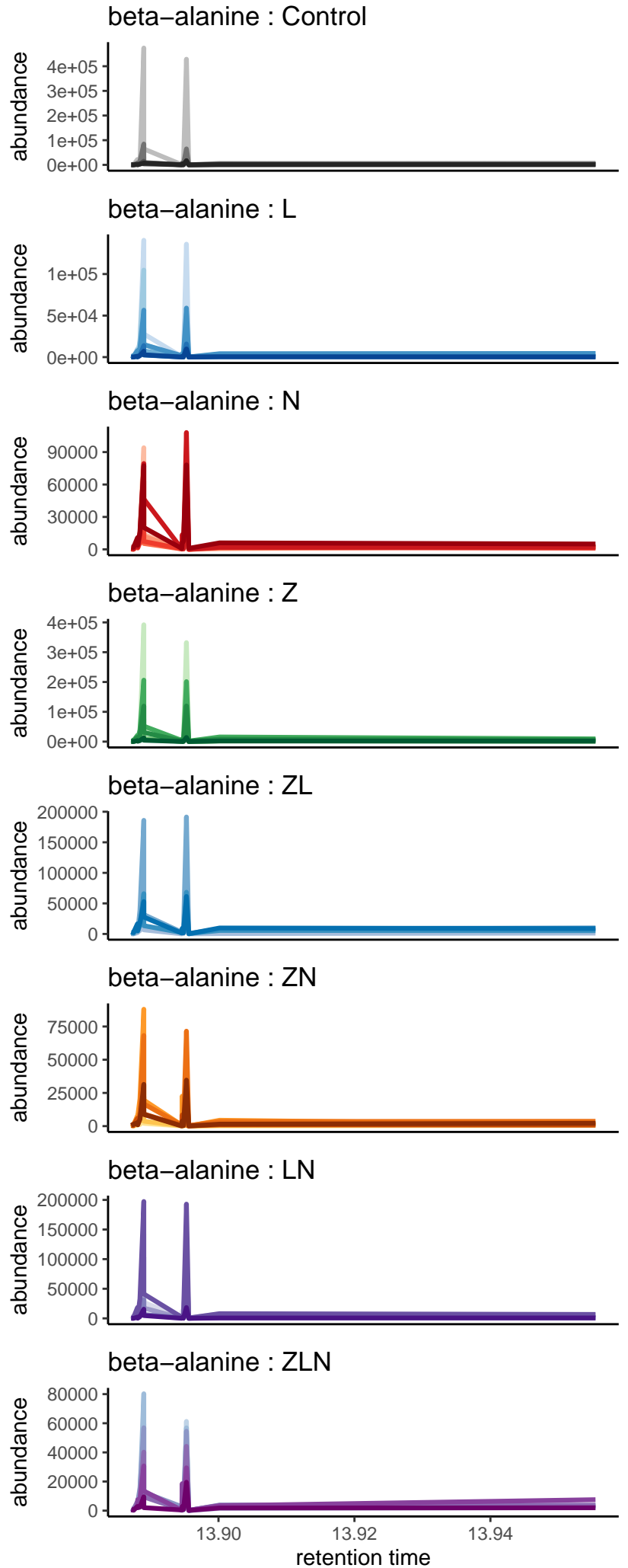
Uracil								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
759	12.42625	0.96	0.02	0.50	0.23	0.24	0.99	0.97
760	12.43190	0.78	0.35	0.36	0.34	0.23	0.02	0.19
761	12.59587	0.27	0.26	0.32	0.20	0.34	0.39	0.05

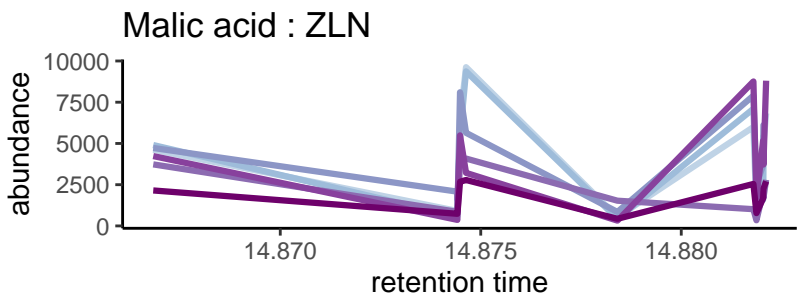
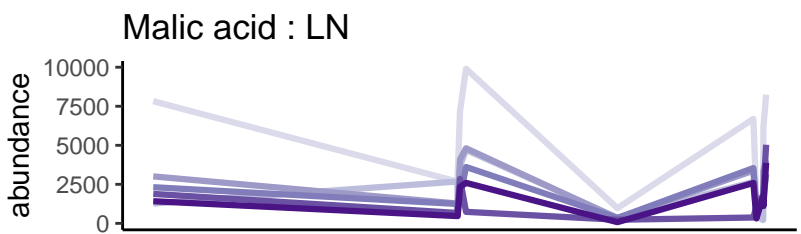
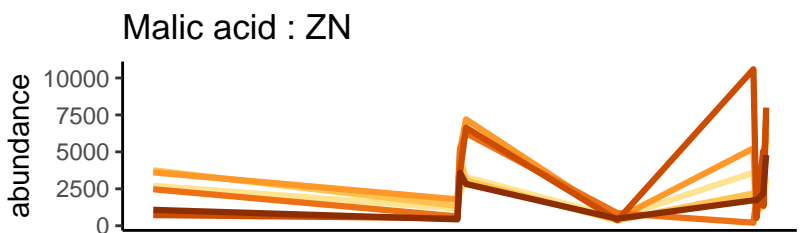
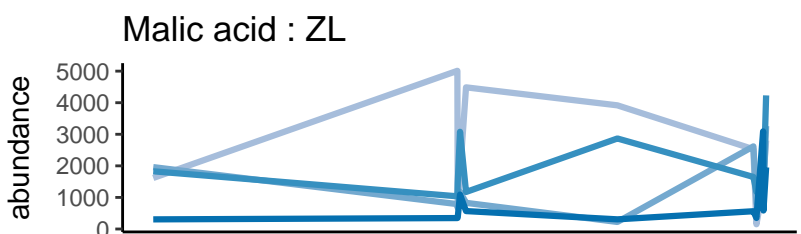
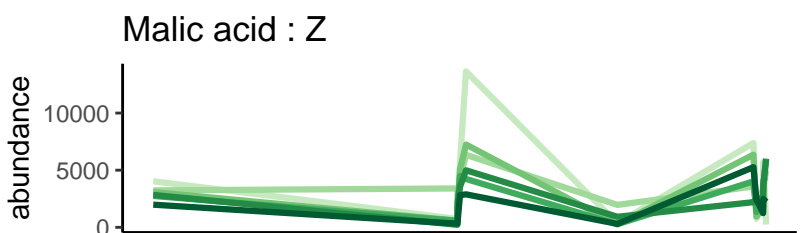
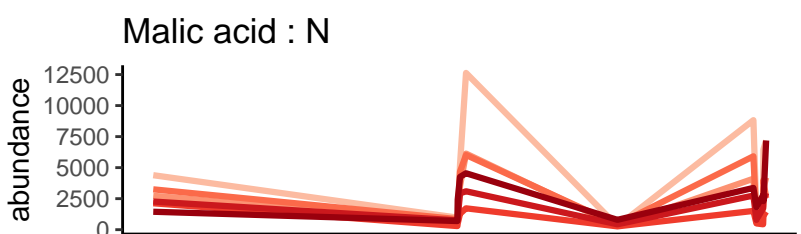
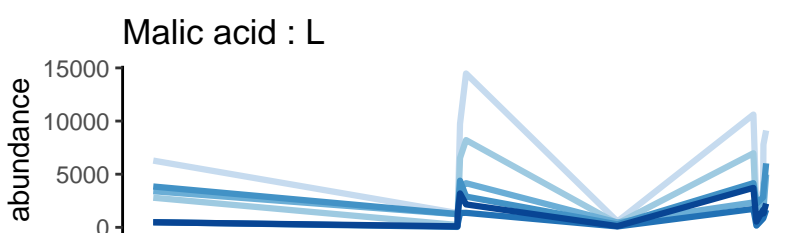
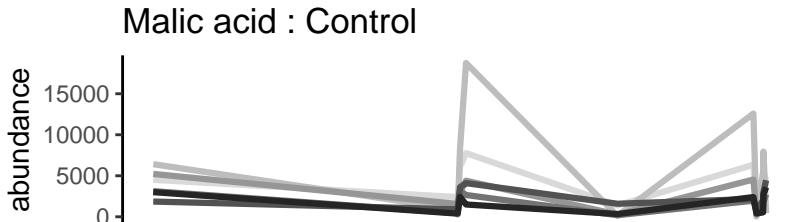
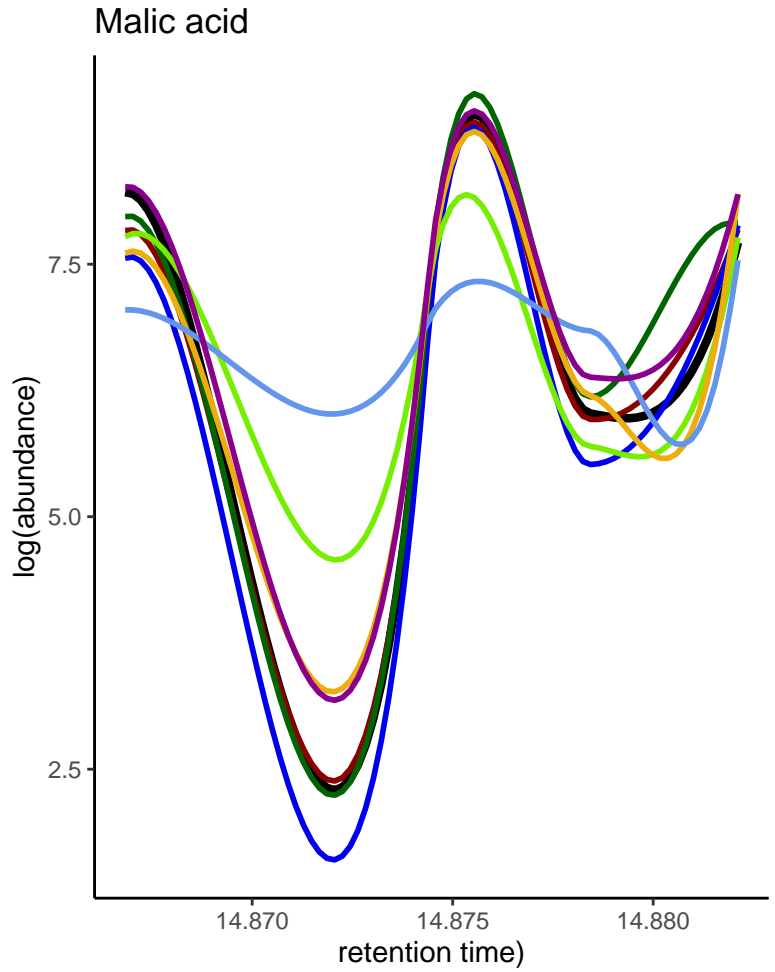






beta-alanine								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
781	13.88717	0.05	0.85	0.69	0.71	0.27	0.56	0.87
782	13.88749	0.05	0.45	0.79	0.75	0.28	0.89	0.98
787	13.88808	0.60	0.99	0.56	0.57	0.02	0.60	0.54
788	13.88812	0.03	0.93	0.74	0.17	0.03	0.60	0.59
789	13.88844	0.45	0.41	0.84	0.89	0.01	0.97	0.60
790	13.88902	0.79	0.88	0.84	0.74	0.03	0.82	0.59
791	13.88902	0.61	0.95	0.71	0.97	0.03	0.99	0.71
798	13.89460	0.16	0.27	0.35	0.90	0.01	0.53	0.80
801	13.89467	0.04	0.43	0.22	0.10	0.35	0.86	0.13
804	13.89484	0.48	0.40	0.52	0.51	0.05	0.85	0.61
806	13.89528	0.69	0.66	0.38	0.57	0.01	0.38	0.22
809	13.89563	0.16	0.56	0.03	1.00	0.22	0.31	0.32
819	13.90016	0.10	0.49	0.58	0.74	0.03	0.27	0.76
828	13.95546	0.09	0.90	0.37	0.04	0.06	0.36	0.82

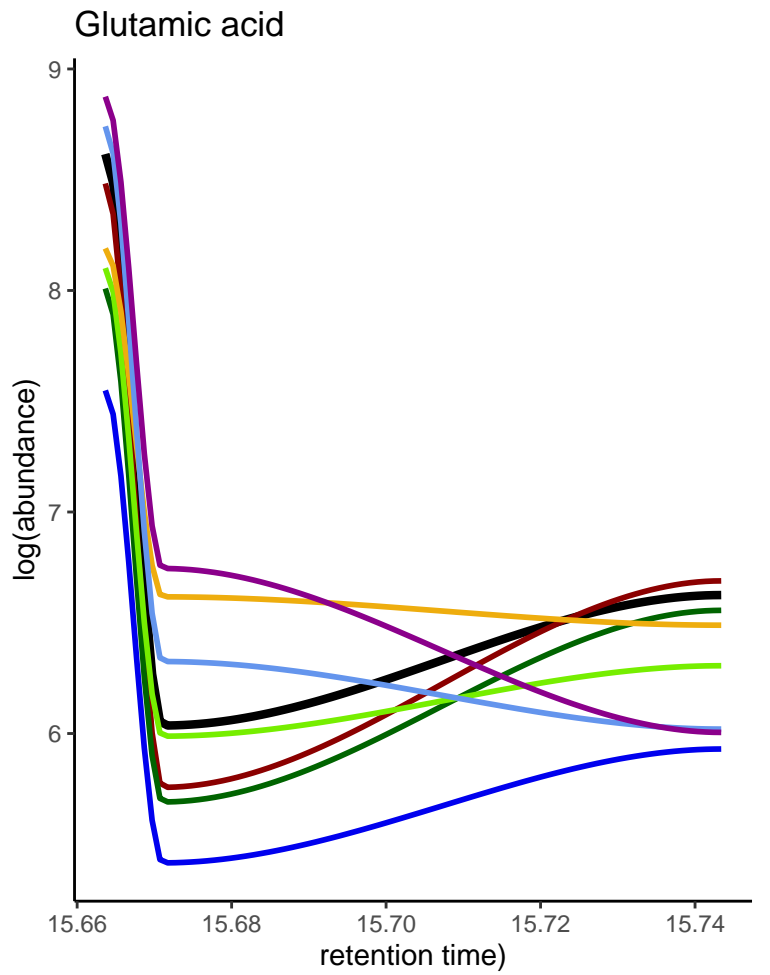




colour

c_mean	ln_mean	z_mean	zln_me
l_mean	n_mean	zl_mean	zn_me

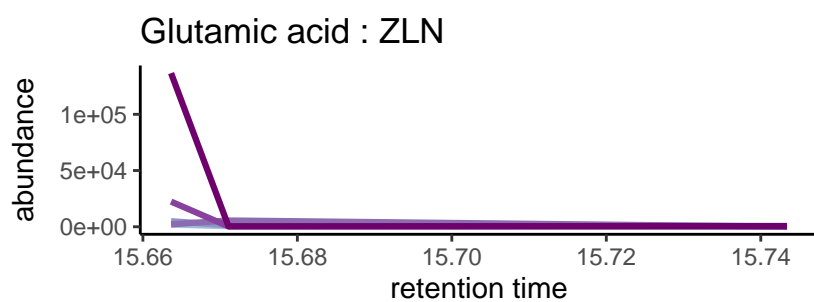
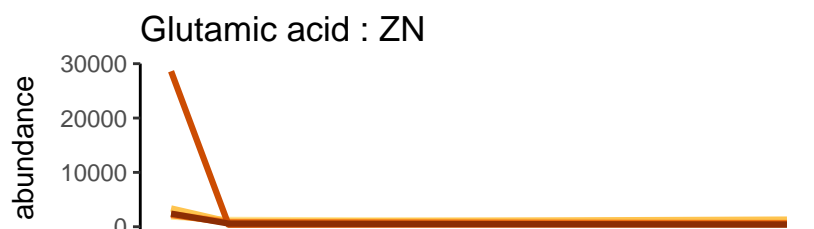
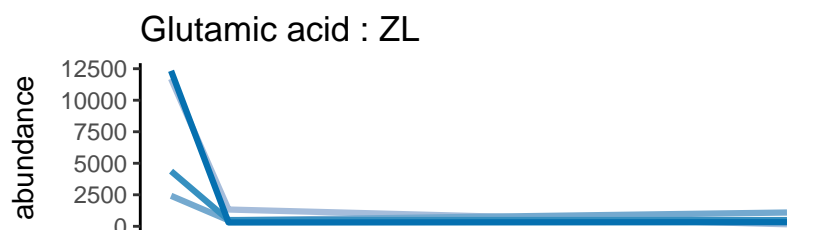
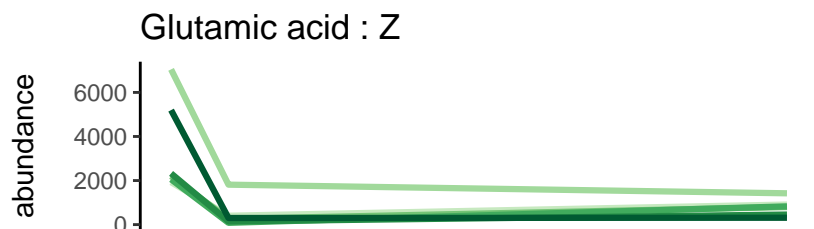
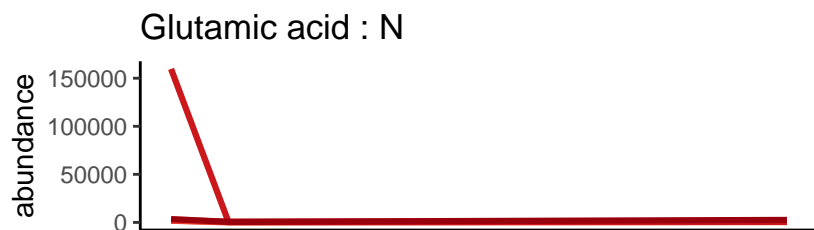
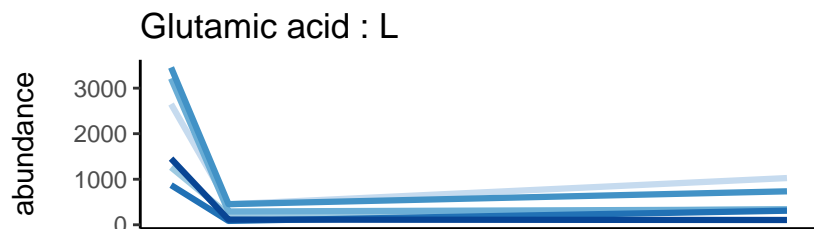
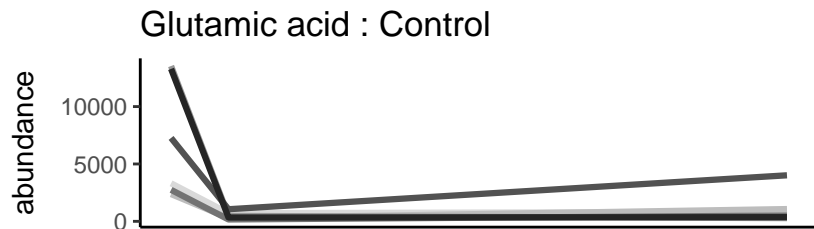
Malic acid								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
847	14.86683	0.75	0.31	0.51	0.38	0.27	0.01	0.20
859	14.87442	0.63	0.94	0.30	0.52	0.40	0.31	0.04
860	14.87443	0.04	0.98	0.13	0.31	0.87	0.63	0.07
862	14.87448	0.97	0.41	0.46	0.39	0.02	0.05	0.03
866	14.87463	0.82	0.07	0.40	0.44	0.20	0.12	0.04
869	14.87841	0.02	0.91	0.87	0.12	0.60	0.84	0.46
878	14.88180	0.62	0.55	0.70	0.81	0.46	0.29	0.04
883	14.88188	0.07	0.78	0.27	0.15	0.63	0.71	0.03
890	14.88204	0.03	0.52	0.72	0.87	0.15	0.71	0.30
891	14.88205	0.99	0.09	0.65	0.92	0.21	0.02	0.03
893	14.88212	0.55	0.37	0.05	0.39	0.34	0.50	0.86

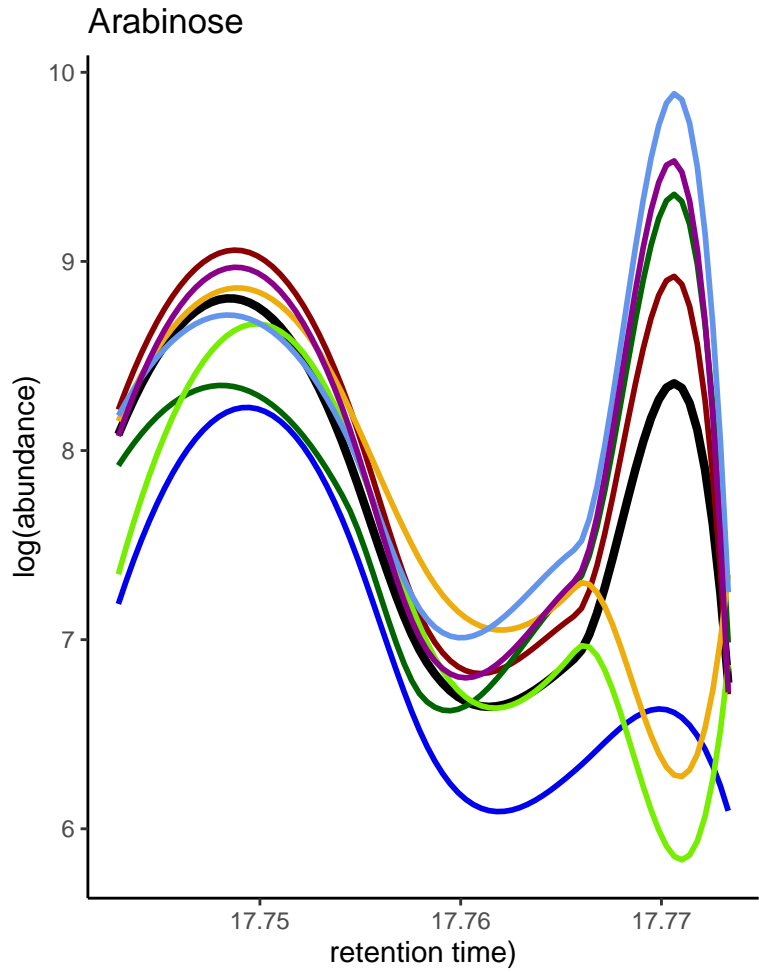


colour

c_mean	ln_mean	z_mean	zln_mean
l_mean	n_mean	zl_mean	zn_mean

Glutamic acid								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
946	15.66365	0.46	0.91	0.49	0.03	0.94	0.60	0.57
949	15.67108	0.04	0.70	0.10	0.26	0.28	0.68	0.19
953	15.74339	0.65	0.02	0.66	0.97	0.55	0.66	0.77

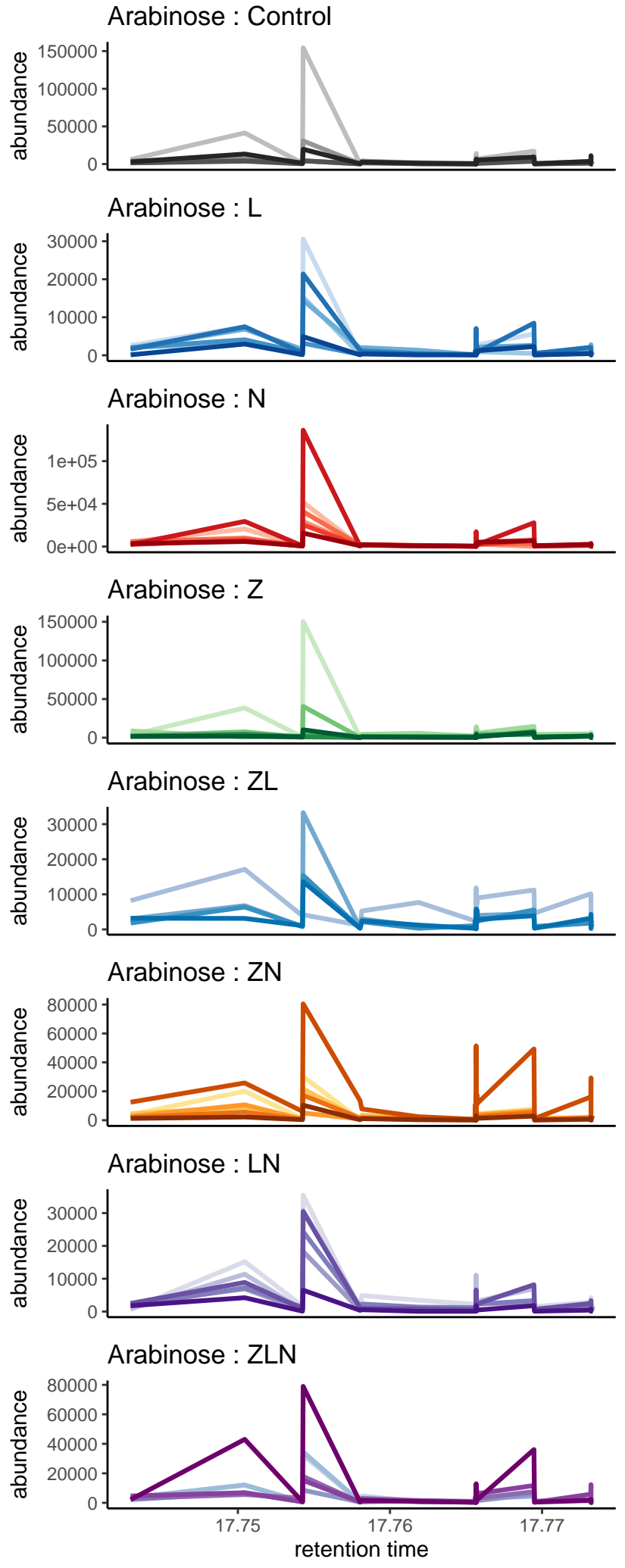




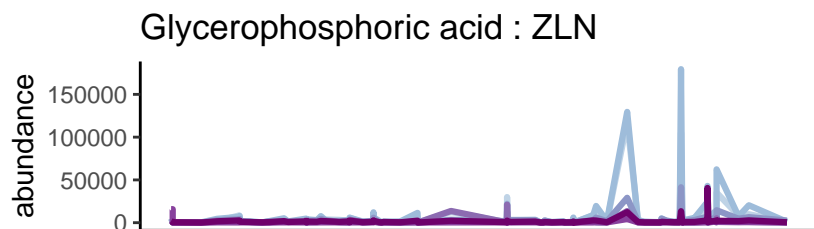
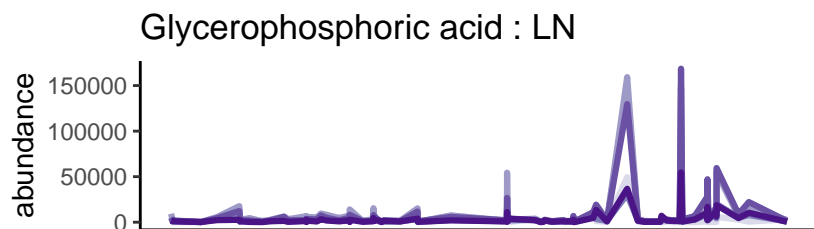
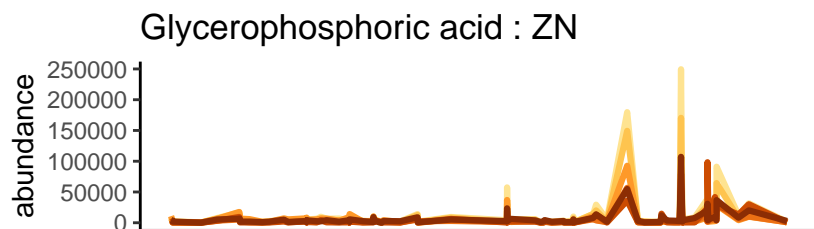
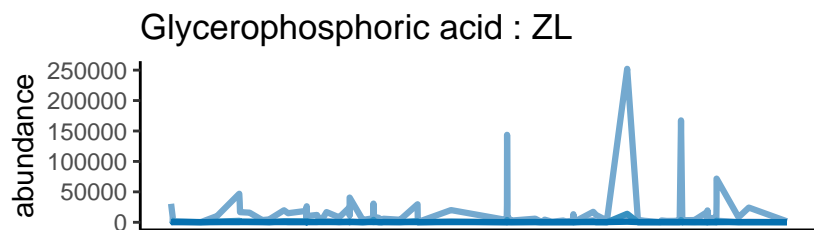
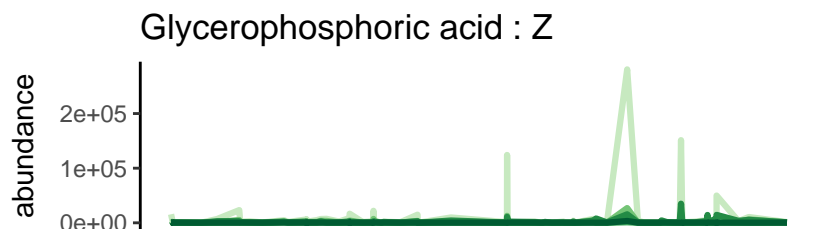
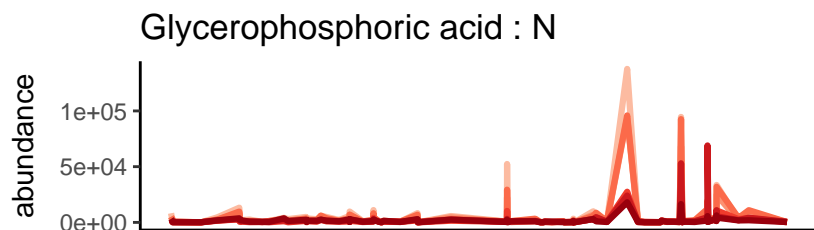
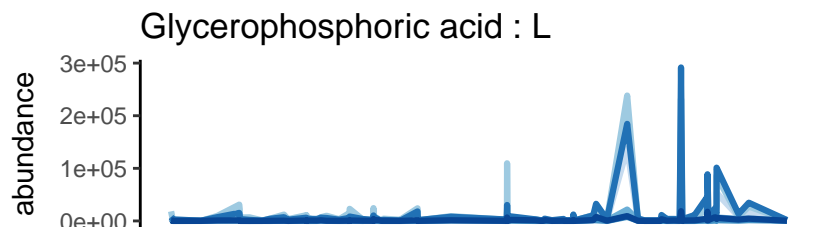
colour

— c\_mean    — ln\_mean    — z\_mean    — zln\_me  
— l\_mean    — n\_mean    — zl\_mean    — zn\_me

Arabinose							
X	retention_time	z_value	l_value	n_value	zl_value	zn_value	ln_value
983	17.74295	0.06	0.04	0.53	0.02	0.81	0.70
985	17.75045	0.70	0.82	0.04	0.11	0.55	0.98
988	17.75425	0.04	0.73	0.80	0.62	0.50	0.85
989	17.75429	0.48	0.66	0.04	0.25	0.86	0.93
990	17.75803	0.94	0.98	0.01	0.59	0.95	0.28
994	17.75815	0.09	0.16	0.84	0.05	0.88	0.72
997	17.76189	0.03	0.20	0.96	0.20	0.33	0.66
999	17.76562	0.01	0.55	0.55	0.23	0.01	0.07
1000	17.76562	0.01	0.76	0.93	0.45	0.38	0.90
1002	17.76563	0.02	0.86	0.64	0.93	0.24	0.78
1006	17.76565	0.02	0.13	0.73	0.23	0.38	0.22
1008	17.76566	0.02	0.58	0.29	0.41	0.04	1.00
1010	17.76567	0.01	0.25	0.29	0.07	0.19	0.96
1011	17.76567	0.04	0.97	0.15	0.78	0.30	0.66
1012	17.76567	0.16	0.34	0.05	0.84	0.78	0.93
1014	17.76568	0.04	0.05	0.27	0.09	0.29	0.45
1015	17.76568	0.04	0.99	0.33	0.14	0.19	0.96
1018	17.76573	0.00	0.48	0.68	0.04	0.75	0.43
1019	17.76575	0.04	0.17	0.26	0.11	0.48	0.51
1020	17.76947	0.04	0.32	0.73	0.45	0.79	0.22
1021	17.76950	0.02	0.34	0.64	0.46	0.05	0.63
1022	17.77320	0.02	0.18	0.67	0.10	0.28	0.73
1023	17.77322	0.01	0.05	0.89	0.09	0.74	0.53
1024	17.77322	0.04	0.74	0.60	0.36	0.04	0.80
1026	17.77332	0.16	0.37	0.26	0.31	0.52	0.93



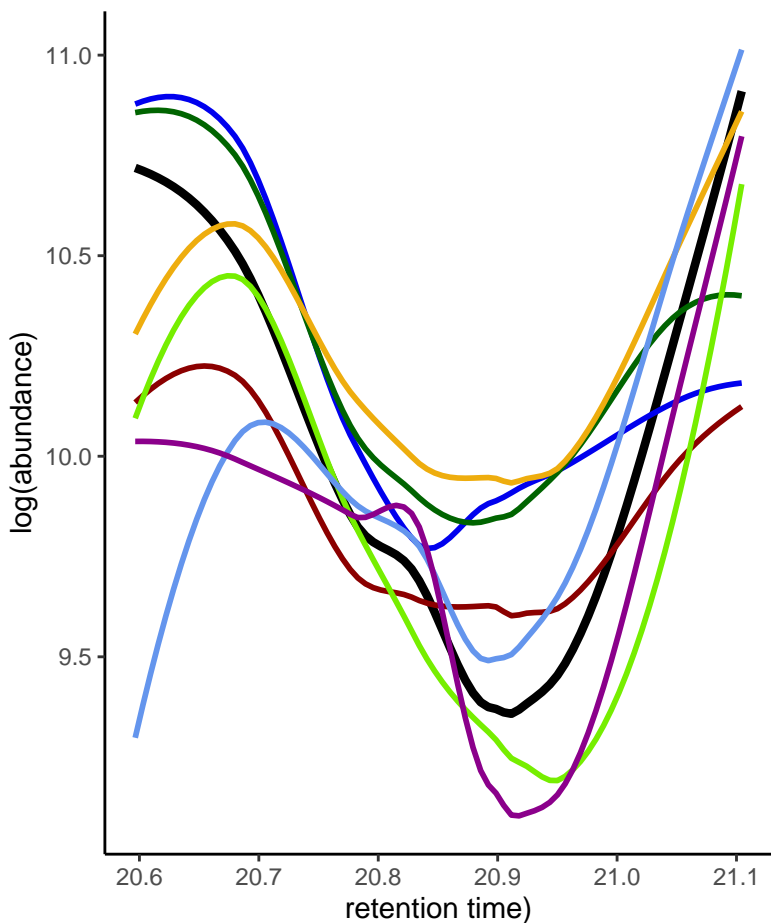
Glycerophosphoric acid : Control



■ c\_mean    ■ ln\_mean    ■ z\_mean    ■ zln\_mean  
■ l\_mean    ■ n\_mean    ■ zl\_mean    ■ zn\_mean

[illegible]

# Glucose



colour

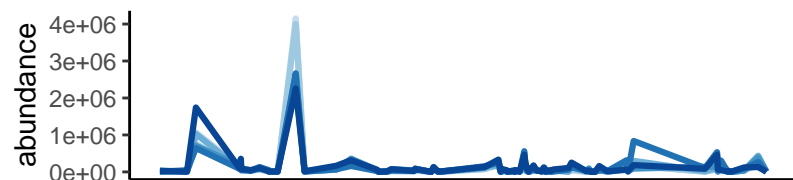
c\_mean ln\_mean z\_mean zln\_me  
l\_mean n\_mean zl\_mean zn\_me

Glucose									
X	retention_time	zln_mean	ln_mean	z_mean	zl_mean	zn_mean	zn_mean	zn_mean	zn_mean
1225	20.6846	0.90	0.34	0.04	0.42	0.11	0.02	0.01	0.01
1227	20.6930	0.13	0.02	0.00	0.15	0.02	0.01	0.01	0.01
1228	20.6950	0.75	0.05	0.70	0.05	0.75	0.05	0.05	0.05
1229	20.6950	0.34	0.14	0.06	0.02	0.02	0.02	0.02	0.02
1227	20.6945	0.77	0.70	0.63	0.05	0.04	0.00	0.00	0.00
1229	20.6943	0.11	0.00	0.13	0.02	0.12	0.00	0.00	0.00
1241	20.6724	0.11	0.17	0.11	0.01	0.00	0.00	0.00	0.00
1242	20.6723	0.94	0.84	0.02	0.05	0.00	0.75	0.00	0.00
1244	20.6707	0.35	0.19	0.09	0.03	0.01	0.00	0.00	0.00
1252	20.6870	0.75	0.30	0.00	0.04	0.03	0.41	0.00	0.00
1253	20.6870	0.13	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1254	20.6870	0.86	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1264	20.6853	0.90	0.40	0.72	0.02	0.70	0.00	0.00	0.00
1270	20.7022	0.76	0.31	0.03	0.00	0.00	0.00	0.00	0.00
1272	20.7161	0.97	0.52	0.30	0.04	0.04	0.42	0.00	0.00
1277	20.7446	0.16	0.40	0.12	0.01	0.00	0.00	0.00	0.00
1281	20.7500	0.20	0.20	0.50	0.01	0.00	0.01	0.00	0.00
1284	20.7363	0.92	0.37	0.97	0.00	0.00	0.00	0.00	0.00
1285	20.7358	0.40	0.31	0.26	0.03	0.04	0.00	0.00	0.00
1288	20.7075	0.03	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1296	20.7036	0.86	0.00	0.70	0.00	0.00	0.00	0.00	0.00
1294	20.6895	0.46	0.52	0.02	0.00	0.00	0.00	0.00	0.00
1295	20.6925	0.86	0.31	0.25	0.14	0.70	0.04	0.00	0.00
1296	20.6910	0.91	0.44	0.01	0.14	0.40	0.00	0.00	0.00
1298	20.6826	0.32	0.37	0.30	0.05	0.04	0.00	0.00	0.00
1300	20.6760	0.40	0.45	0.01	0.00	0.07	0.04	0.00	0.00
1305	20.6977	0.03	0.20	0.24	0.20	0.10	0.00	0.00	0.00
1307	20.6868	0.03	0.00	0.01	0.00	0.00	0.00	0.00	0.00
1308	20.6790	0.46	0.40	0.71	0.02	0.44	0.00	0.00	0.00
1310	20.6810	0.12	0.70	0.00	0.04	0.00	0.00	0.00	0.00
1318	20.6870	0.12	0.00	0.30	0.00	0.00	0.00	0.00	0.00
1348	20.6877	0.16	0.54	0.03	0.04	0.00	0.00	0.00	0.00
1349	20.6877	0.16	0.30	0.04	0.06	0.10	0.00	0.00	0.00
1370	20.6901	0.02	0.30	0.74	0.23	0.00	0.00	0.00	0.00
1380	20.6975	0.05	0.00	0.00	0.01	0.00	0.00	0.00	0.00
1382	20.6947	0.00	0.00	0.01	0.04	0.00	0.00	0.00	0.00
1388	20.6870	0.11	0.75	0.05	0.24	0.00	0.00	0.00	0.00
1389	20.6820	0.12	0.30	0.04	0.11	0.00	0.04	0.00	0.00
1398	20.6917	0.05	0.22	0.60	0.04	0.00	0.00	0.00	0.00
1401	20.6946	0.40	0.34	0.00	0.00	0.00	0.00	0.00	0.00
1402	20.6907	0.41	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1404	20.6924	0.97	0.71	0.05	0.04	0.25	0.00	0.00	0.00
1407	20.6944	0.42	0.17	0.40	0.00	0.73	0.04	0.00	0.00
1416	20.6910	0.75	0.00	0.04	0.02	0.14	0.00	0.00	0.00
1420	20.6940	0.02	0.34	0.97	0.03	0.00	0.00	0.00	0.00
1422	20.6772	0.20	0.27	0.40	0.04	0.00	0.00	0.00	0.00
1424	20.6924	0.05	0.00	0.24	0.02	0.00	0.00	0.00	0.00
1425	20.6900	0.77	0.02	0.04	0.03	0.12	0.00	0.00	0.00
1428	20.6922	0.37	0.06	0.62	0.00	0.00	0.00	0.00	0.00
1430	20.6920	0.71	0.17	0.00	0.00	0.00	0.00	0.00	0.00
1435	20.6920	0.10	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1440	20.6908	0.11	0.13	0.97	0.02	0.00	0.00	0.00	0.00
1448	20.6915	0.77	0.20	0.54	0.02	0.10	0.00	0.00	0.00
1453	20.6914	0.77	0.44	0.31	0.00	0.00	0.00	0.00	0.00
1456	20.6910	0.04	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1458	20.6940	0.45	0.00	0.00	0.02	0.00	0.00	0.00	0.00
1460	20.6903	0.12	0.24	0.50	0.00	0.00	0.00	0.00	0.00
1464	20.6946	0.04	0.35	0.00	0.00	0.00	0.00	0.00	0.00
1465	20.6927	0.24	0.00	0.26	0.00	0.00	0.00	0.00	0.00
1466	20.6974	0.12	0.41	0.00	0.04	0.10	0.00	0.00	0.00
1468	20.6945	0.02	0.00	0.40	0.10	0.00	0.00	0.00	0.00
1470	20.6910	0.75	0.00	0.00	0.12	0.14	0.00	0.00	0.00
1484	21.0200	0.04	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1485	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1486	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1487	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1488	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1489	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1490	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1491	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1492	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1493	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1494	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1495	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1496	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1497	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1498	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1499	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1500	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1501	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1502	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1503	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1504	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1505	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1506	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1507	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1508	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1509	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1510	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
1511	21.0200	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00

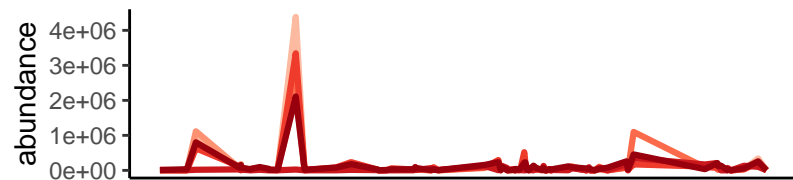
## Glucose : Control



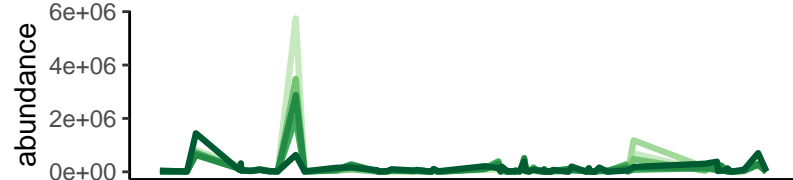
## Glucose : L



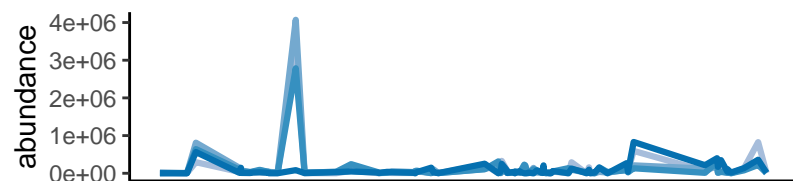
## Glucose : N



## Glucose : Z



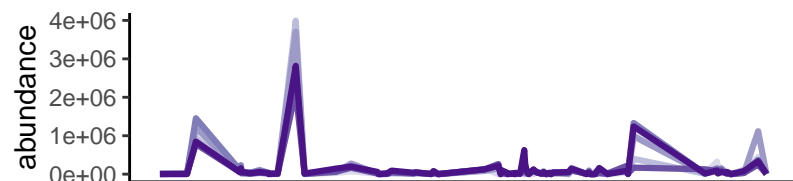
## Glucose : ZL



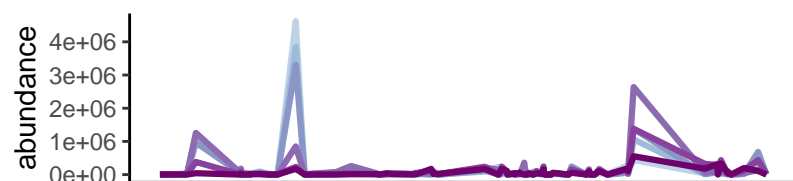
## Glucose : ZN



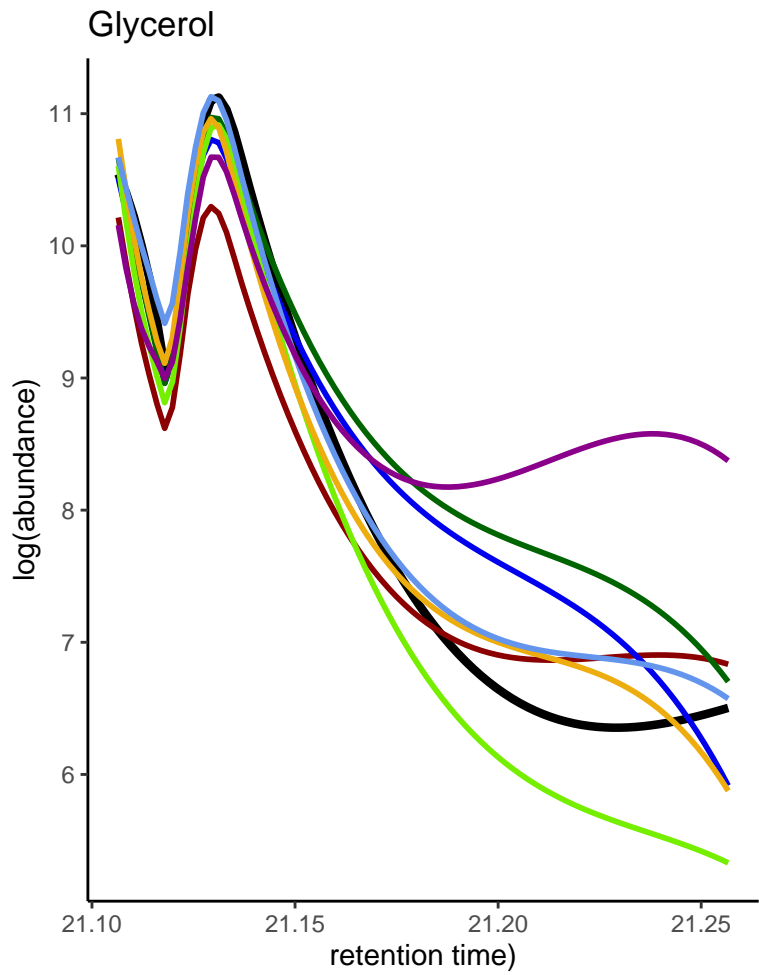
## Glucose : LN



## Glucose : ZLN



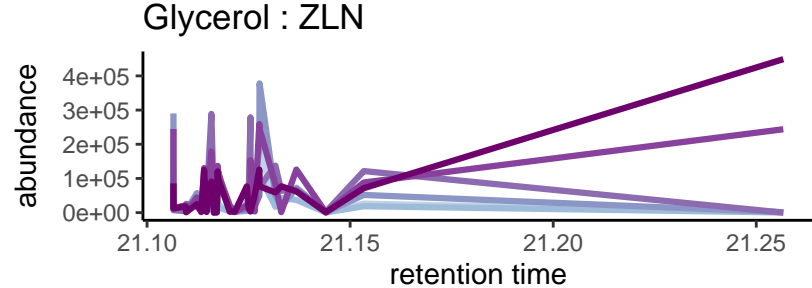
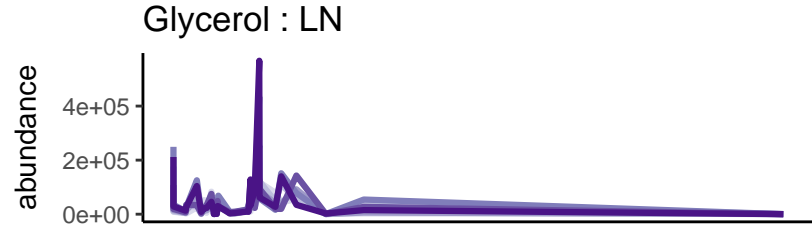
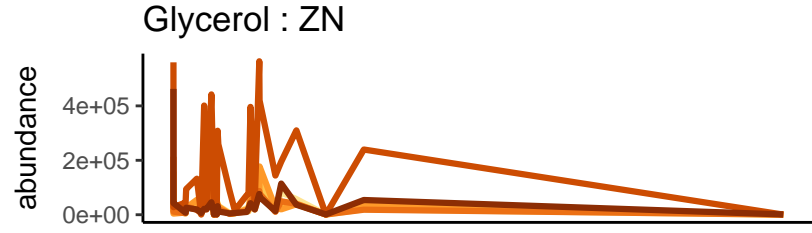
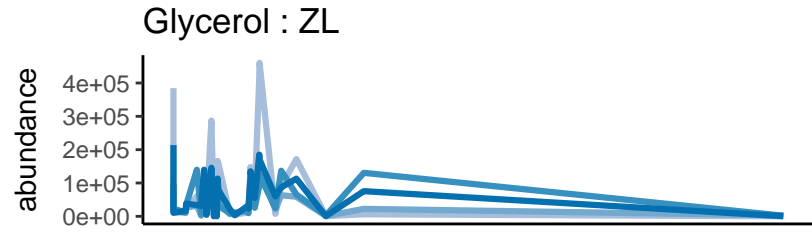
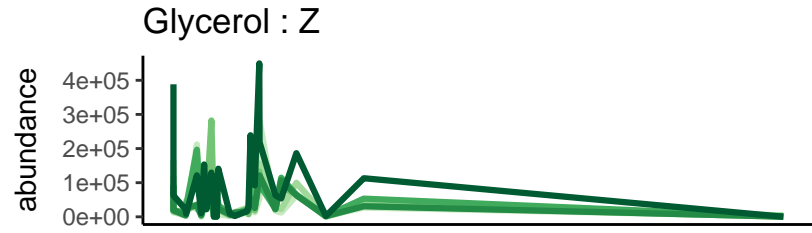
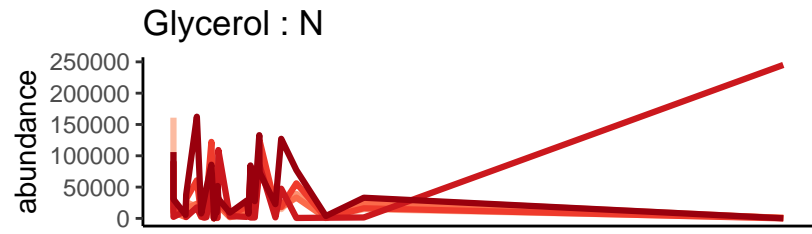
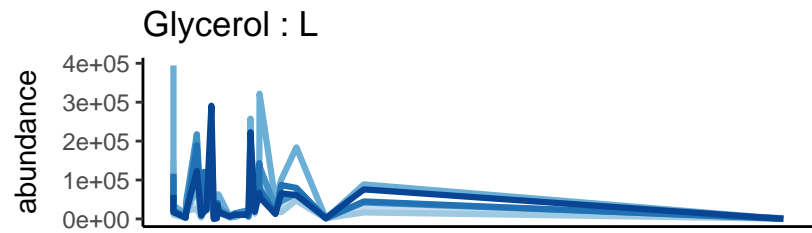
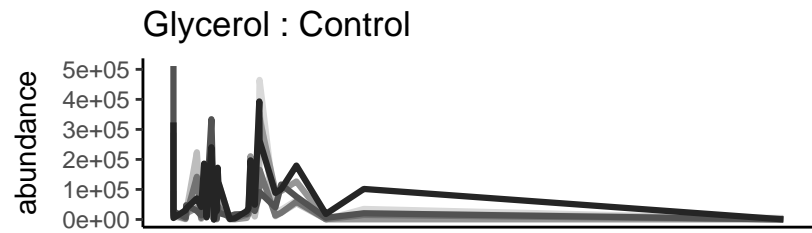
retention time



colour

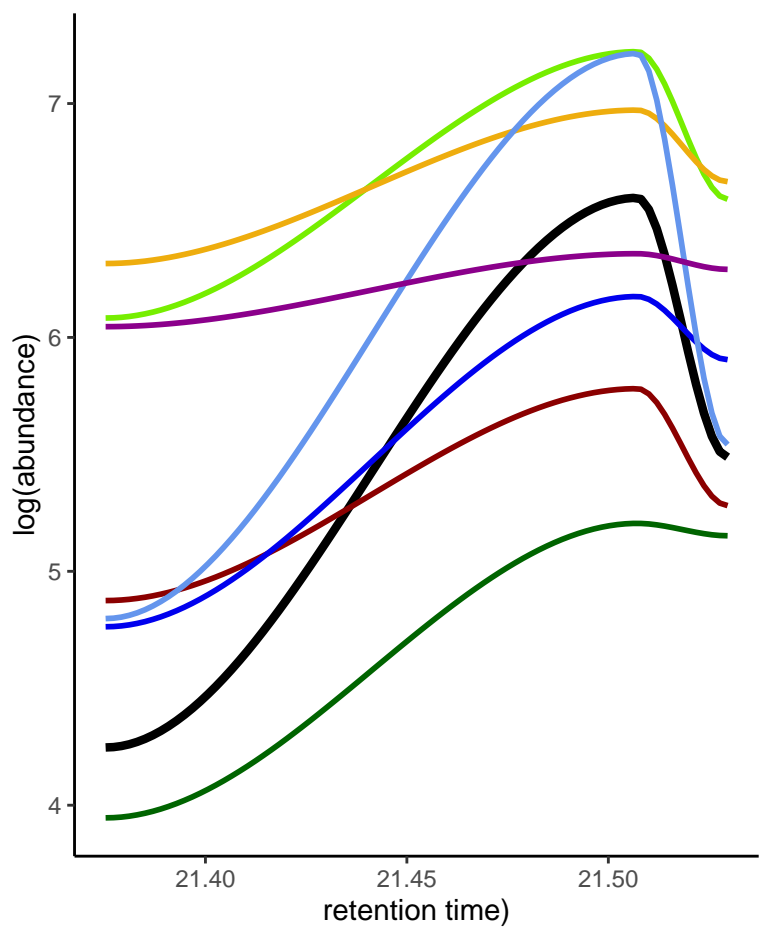
— c\_mean    — ln\_mean    — z\_mean    — zln\_me  
— l\_mean    — n\_mean    — zl\_mean    — zn\_me

Glycerol								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
1523	21.10650	0.56	0.56	0.57	0.70	0.25	0.61	0.03
1524	21.10650	0.63	0.71	0.26	0.04	0.71	0.75	0.94
1531	21.10949	0.05	0.93	0.71	0.90	0.53	1.00	0.15
1533	21.10963	0.41	0.07	0.16	0.10	0.72	0.03	0.01
1542	21.11227	0.31	0.89	0.01	0.31	1.00	0.65	0.90
1543	21.11333	0.38	0.75	0.03	0.48	0.45	0.58	0.81
1546	21.11406	0.13	0.93	0.05	0.29	0.04	0.24	0.13
1547	21.11463	0.91	0.24	0.17	0.02	0.66	0.28	0.86
1554	21.11584	0.32	0.40	0.01	0.15	0.02	0.92	0.91
1556	21.11643	0.15	0.28	0.92	0.73	0.57	0.48	0.02
1558	21.11647	0.03	0.52	0.75	0.29	0.44	0.38	0.17
1565	21.11692	0.95	0.49	0.02	0.26	0.69	0.32	0.28
1567	21.11715	0.05	0.33	0.36	0.98	0.44	0.63	0.49
1569	21.11738	0.04	0.38	0.18	0.42	0.22	0.39	0.24
1570	21.11755	0.49	0.55	0.88	0.02	0.77	0.51	0.19
1587	21.12037	0.03	0.25	0.93	0.46	0.51	0.19	0.78
1589	21.12161	0.49	0.67	0.31	0.11	0.74	0.03	0.40
1598	21.12456	0.05	0.70	0.99	0.60	0.61	0.63	0.82
1599	21.12497	0.01	0.74	0.06	0.55	0.01	0.09	0.00
1602	21.12547	0.47	0.08	0.00	0.38	0.60	0.19	0.32
1604	21.12548	0.50	0.96	0.04	0.49	0.26	0.98	0.44
1605	21.12652	0.53	0.26	0.41	0.36	0.72	0.37	0.00
1610	21.12763	0.81	0.45	0.90	0.91	0.68	0.07	0.02
1611	21.12770	0.19	0.33	0.01	0.07	0.15	0.40	0.24
1623	21.13160	0.32	0.36	0.87	0.87	0.30	0.02	0.56
1627	21.13303	0.88	0.82	0.24	0.22	0.52	0.67	0.03
1631	21.13677	0.25	0.23	0.04	0.54	0.48	0.17	0.22
1641	21.14403	0.55	0.31	0.03	0.97	0.96	0.73	0.38
1645	21.15338	0.02	0.34	0.53	0.42	0.28	0.65	0.29
1649	21.25665	0.19	0.86	0.79	0.04	0.69	0.52	0.14





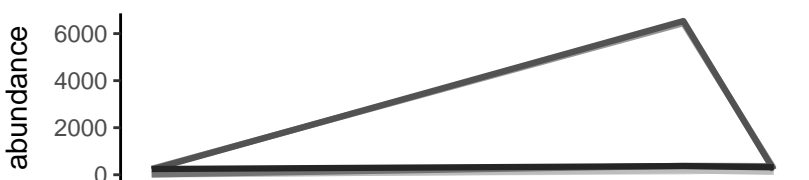
Glucitol/gluconic acid



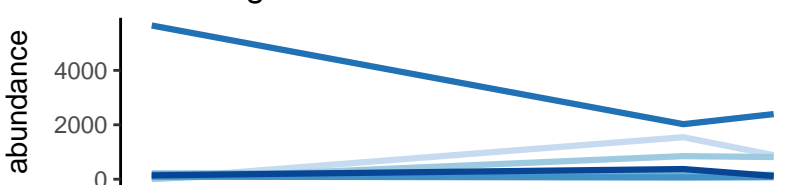
colour    c\_mean    ln\_mean    z\_mean    zln\_mean  
          l\_mean    n\_mean    zl\_mean    zn\_mean

Glucitol/gluconic acid								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
1653	21.37523	0.59	0.30	0.03	0.64	0.47	0.90	0.47
1654	21.50719	0.85	0.18	0.40	0.93	0.56	0.76	0.01
1656	21.52960	0.67	0.11	0.04	0.17	0.13	0.85	0.18

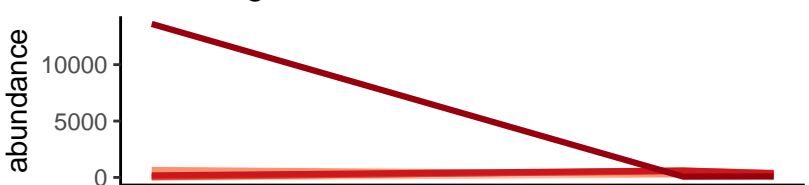
Glucitol/gluconic acid : Control



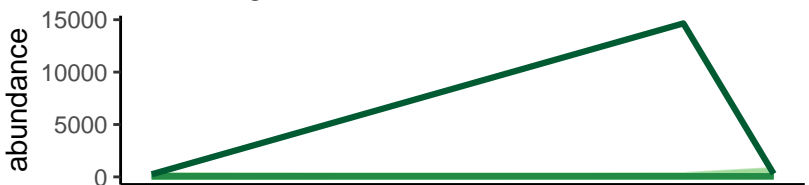
Glucitol/gluconic acid : L



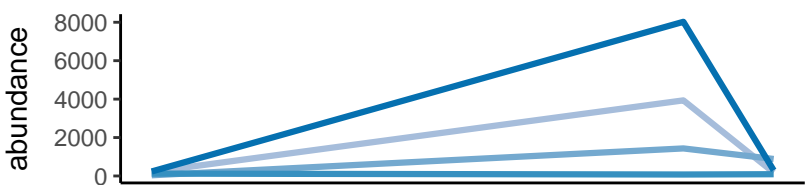
Glucitol/gluconic acid : N



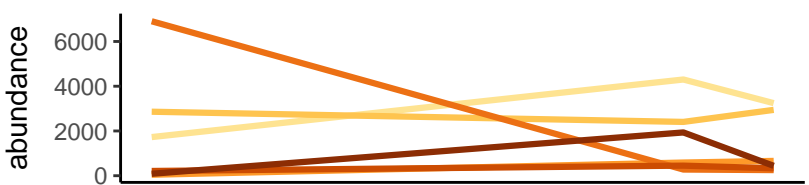
Glucitol/gluconic acid : Z



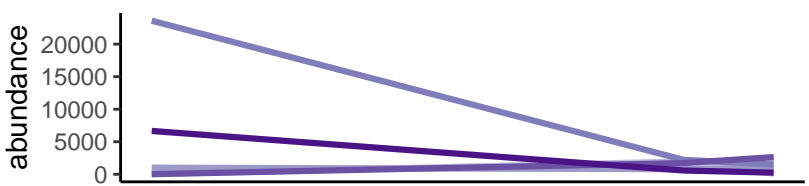
Glucitol/gluconic acid : ZL



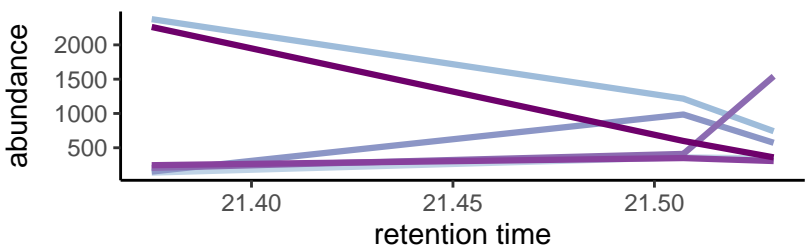
Glucitol/gluconic acid : ZN



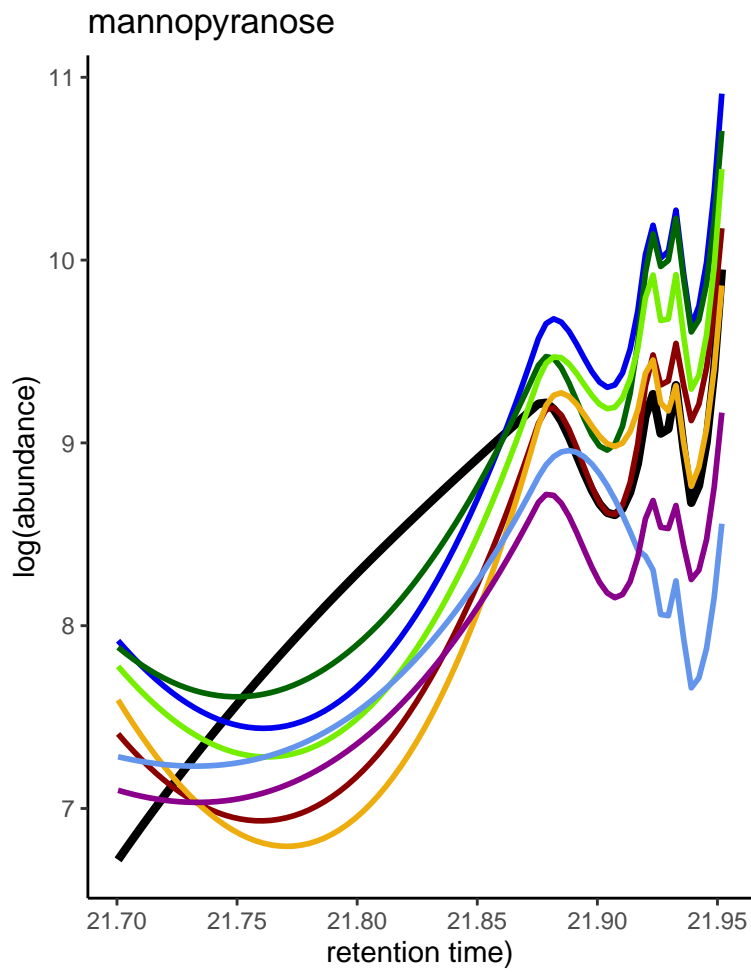
Glucitol/gluconic acid : LN



Glucitol/gluconic acid : ZLN



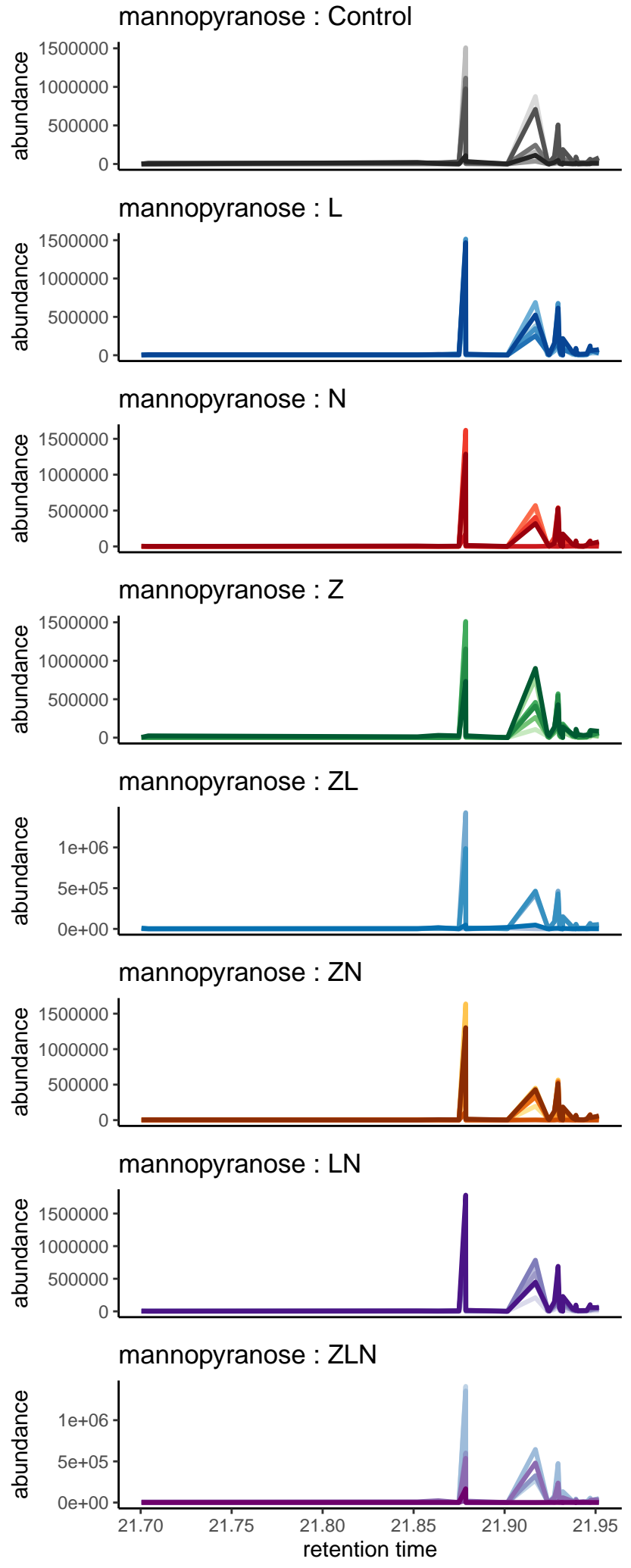


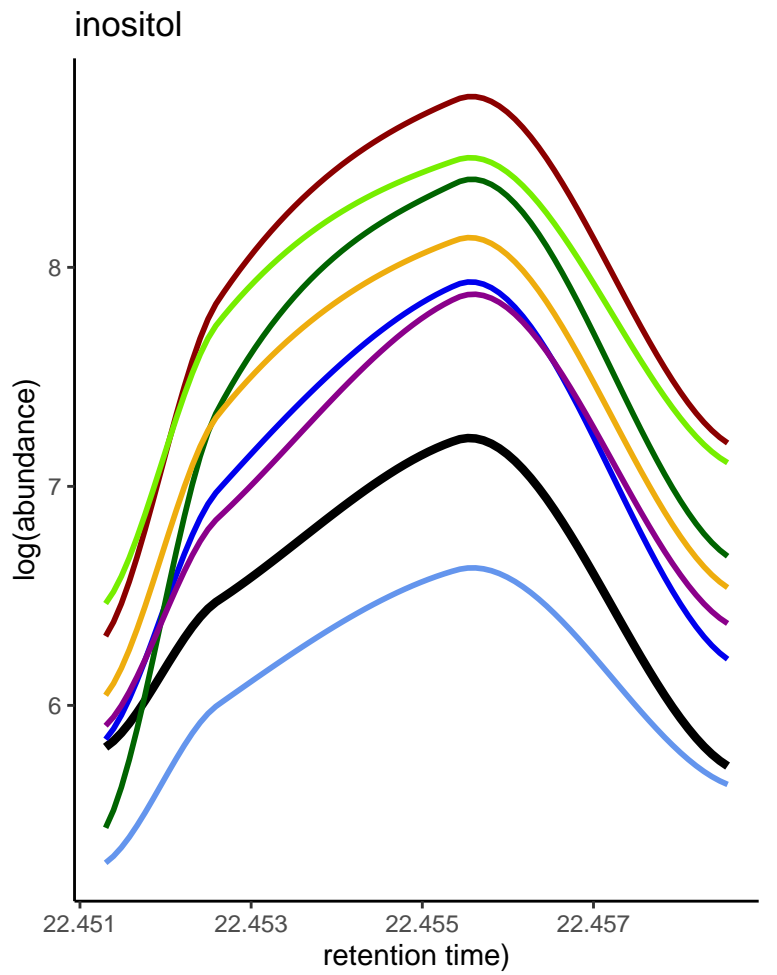


colour

c\_mean    ln\_mean    z\_mean    zn\_me  
 l\_mean    n\_mean    zl\_mean    zn\_me

mannopyranose								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zn_fvalue
1658	21.70046	0.44	0.36	0.98	0.04	0.08	0.18	0.59
1659	21.70418	0.57	0.96	0.87	0.03	0.98	0.92	0.36
1667	21.85210	0.13	0.60	0.03	0.84	0.81	0.13	0.78
1670	21.86356	0.88	0.51	0.04	0.59	0.78	0.90	0.20
1676	21.87493	0.48	0.61	0.69	0.04	0.83	0.87	0.16
1677	21.87864	0.17	0.31	0.62	0.01	0.97	0.92	0.12
1679	21.87870	0.04	0.20	0.34	0.25	0.86	0.33	0.45
1689	21.90160	0.38	0.05	0.47	0.86	0.75	0.04	0.01
1697	21.91695	0.18	0.64	0.42	0.04	0.64	0.67	0.90
1699	21.92356	0.17	0.28	0.79	0.02	0.78	0.71	0.17
1700	21.92377	0.43	0.56	0.82	0.04	0.69	0.93	0.12
1701	21.92430	0.40	0.72	0.89	0.02	0.81	0.97	0.15
1702	21.92440	0.29	0.50	0.70	0.04	0.90	0.91	0.17
1704	21.92447	0.25	0.70	0.87	0.03	0.81	0.82	0.19
1705	21.92453	0.43	0.54	0.89	0.02	0.99	0.61	0.14
1706	21.92739	0.18	0.51	0.61	0.02	0.92	0.49	0.44
1708	21.92942	0.28	0.52	0.60	0.02	0.88	0.73	0.64
1709	21.93017	0.11	0.69	0.82	0.05	0.90	0.77	0.22
1710	21.93093	0.20	0.56	0.57	0.05	0.85	0.38	0.07
1713	21.93165	0.15	0.52	0.75	0.02	0.51	0.96	0.19
1714	21.93175	0.47	0.62	0.94	0.02	0.43	0.87	0.05
1717	21.93193	0.03	0.51	0.29	0.02	0.32	0.83	0.42
1718	21.93195	0.76	0.63	0.03	0.24	0.30	0.56	0.77
1719	21.93200	0.22	0.47	0.81	0.01	0.83	0.70	0.39
1722	21.93742	0.18	0.60	0.82	0.02	0.54	0.87	0.09
1724	21.93913	0.04	0.17	0.59	0.02	0.50	0.67	0.18
1725	21.93916	0.13	0.67	0.93	0.01	0.61	0.78	0.06
1726	21.93919	0.21	0.54	0.97	0.03	0.78	0.86	0.04
1727	21.93920	0.15	0.46	0.66	0.04	0.85	0.80	0.19
1729	21.93925	0.09	0.47	0.61	0.03	0.84	0.39	0.35
1731	21.94025	0.38	0.43	0.65	0.04	0.93	0.92	0.23
1733	21.94335	0.56	0.93	0.43	0.04	0.98	0.57	0.29
1734	21.94523	0.09	0.35	0.93	0.02	0.92	0.57	0.19
1736	21.94704	0.12	0.39	0.65	0.02	0.55	0.44	0.32
1737	21.94725	0.14	0.44	0.52	0.04	0.94	0.45	0.13
1739	21.95182	0.13	0.56	0.81	0.05	0.89	0.52	0.22

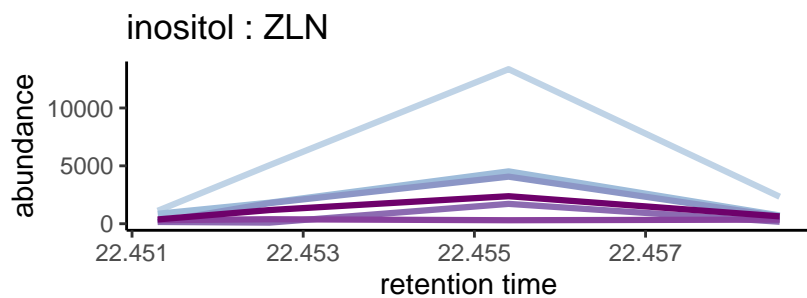
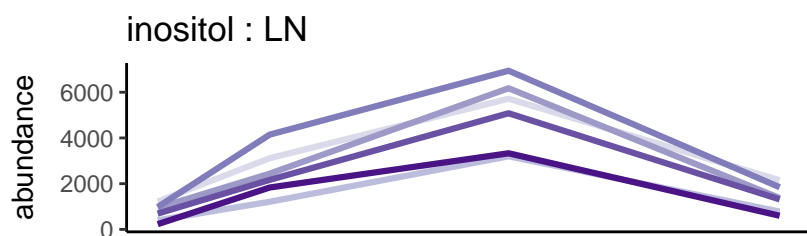
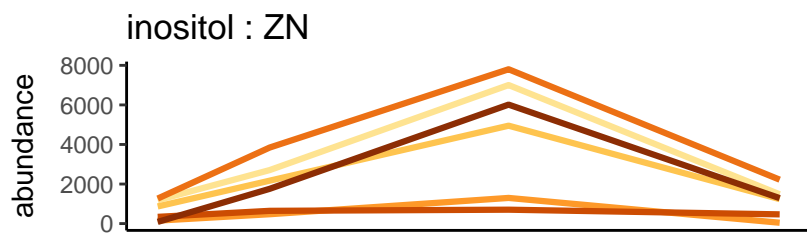
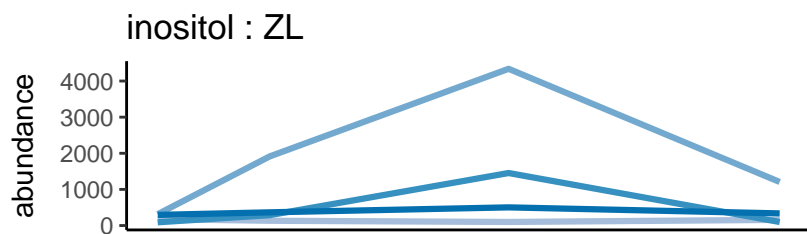
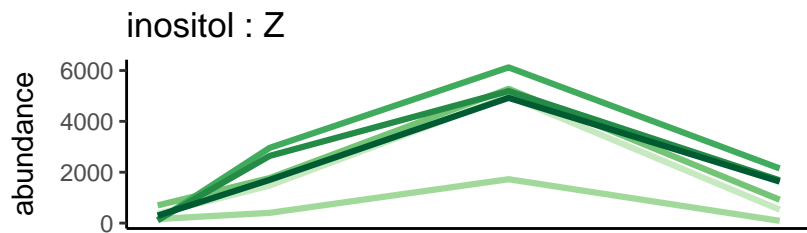
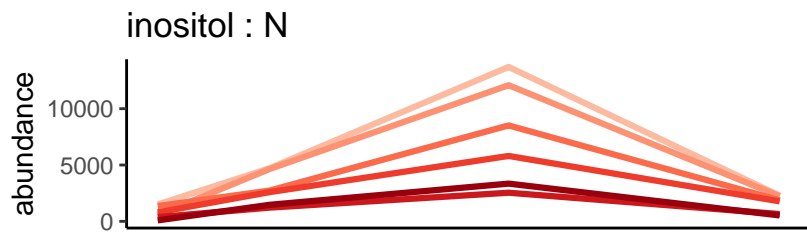
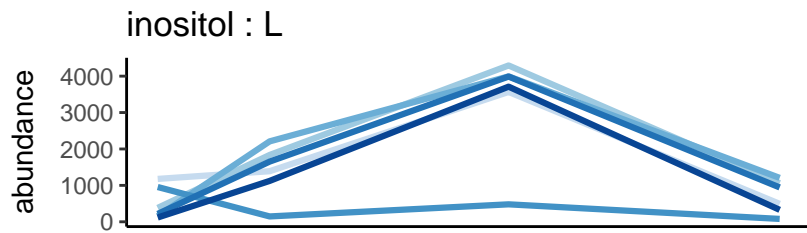
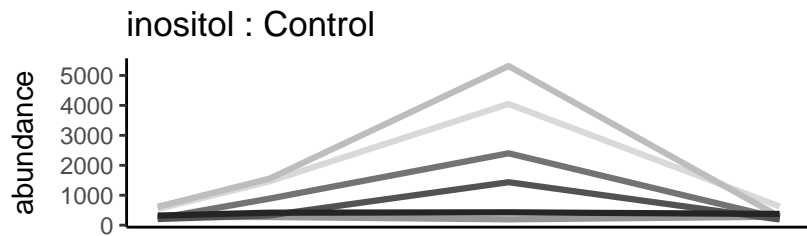


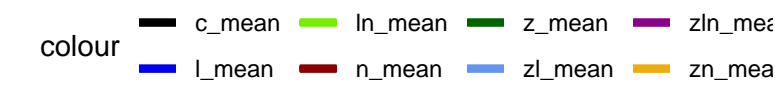
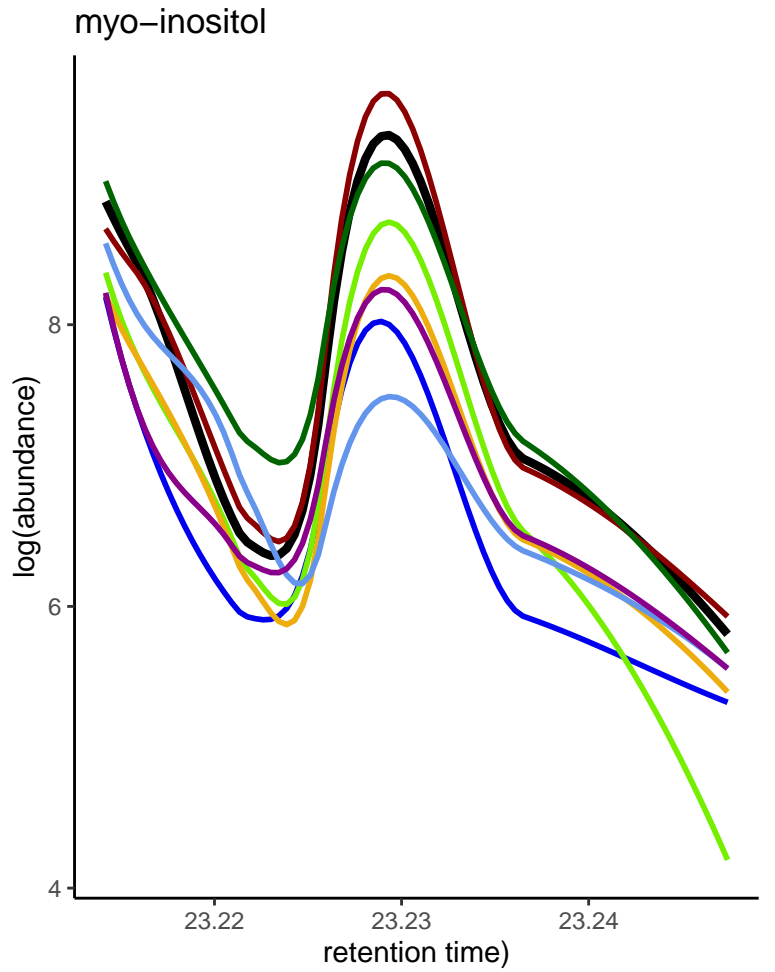


colour

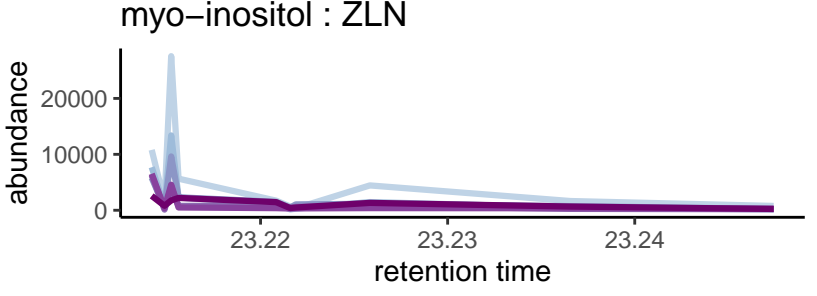
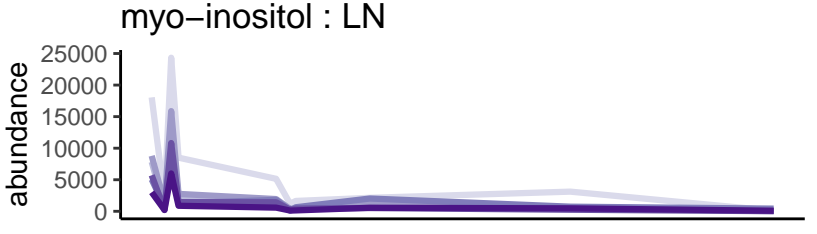
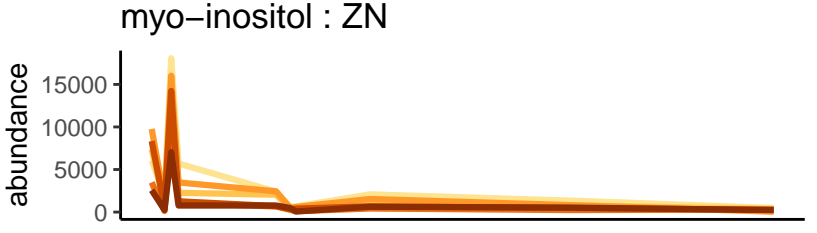
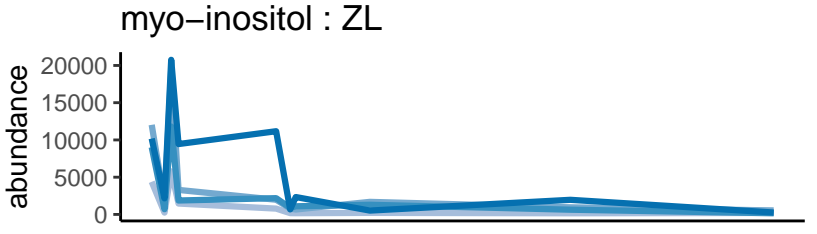
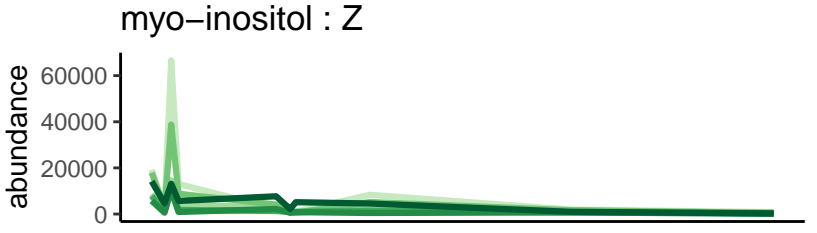
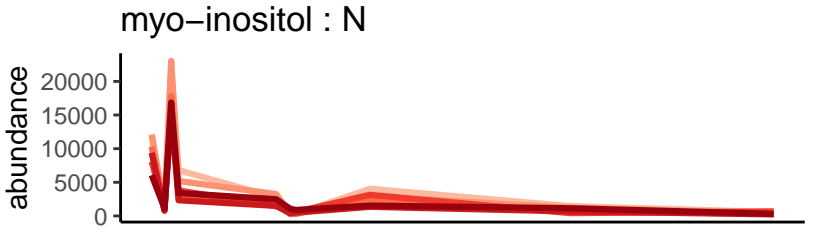
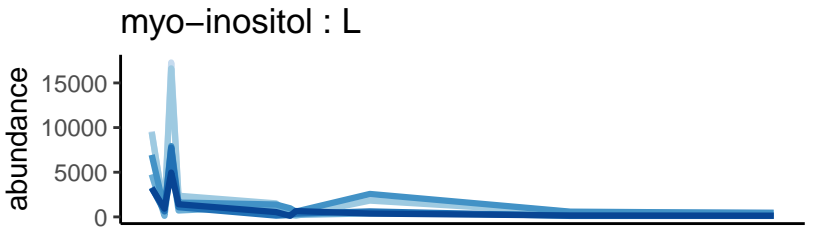
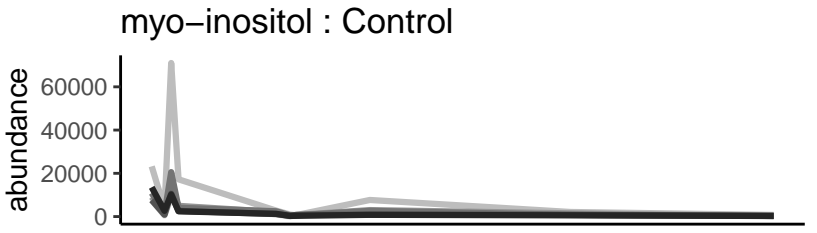
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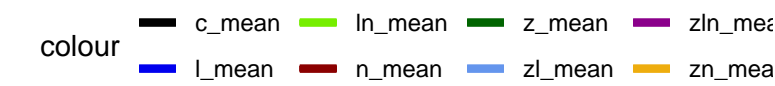
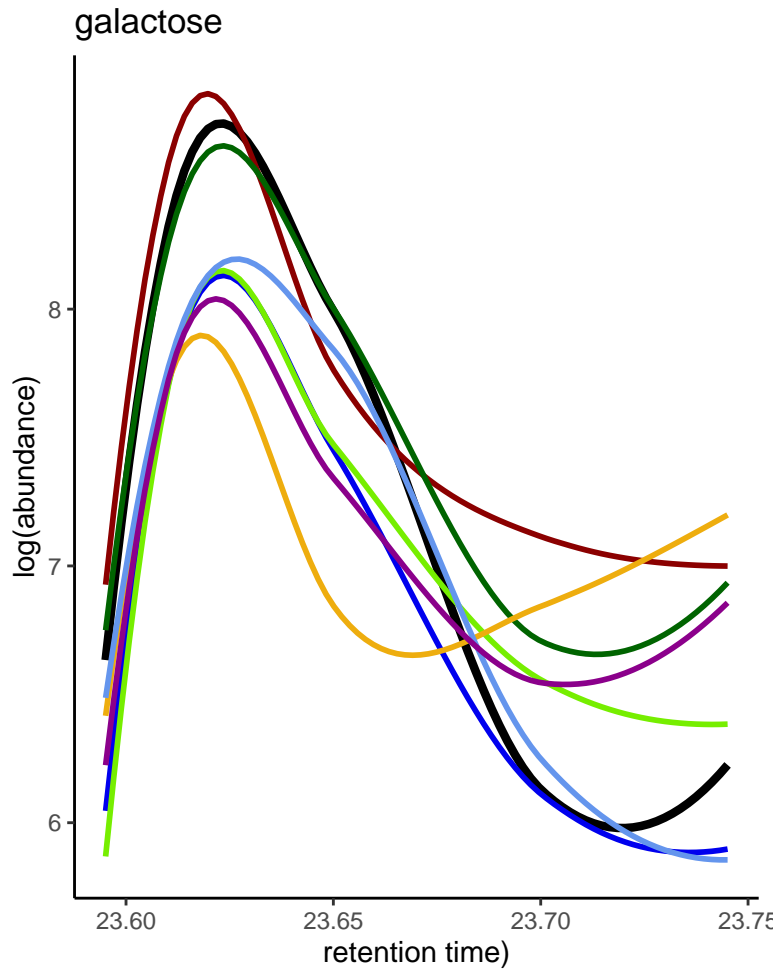
inositol								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
1747	22.45130	0.11	0.98	0.02	0.64	0.92	0.89	0.93
1748	22.45261	0.26	0.28	0.01	0.04	0.19	0.86	0.17
1749	22.45540	0.40	0.29	0.02	0.05	0.27	0.75	0.04
1750	22.45857	0.50	0.60	0.01	0.16	0.10	0.85	0.21



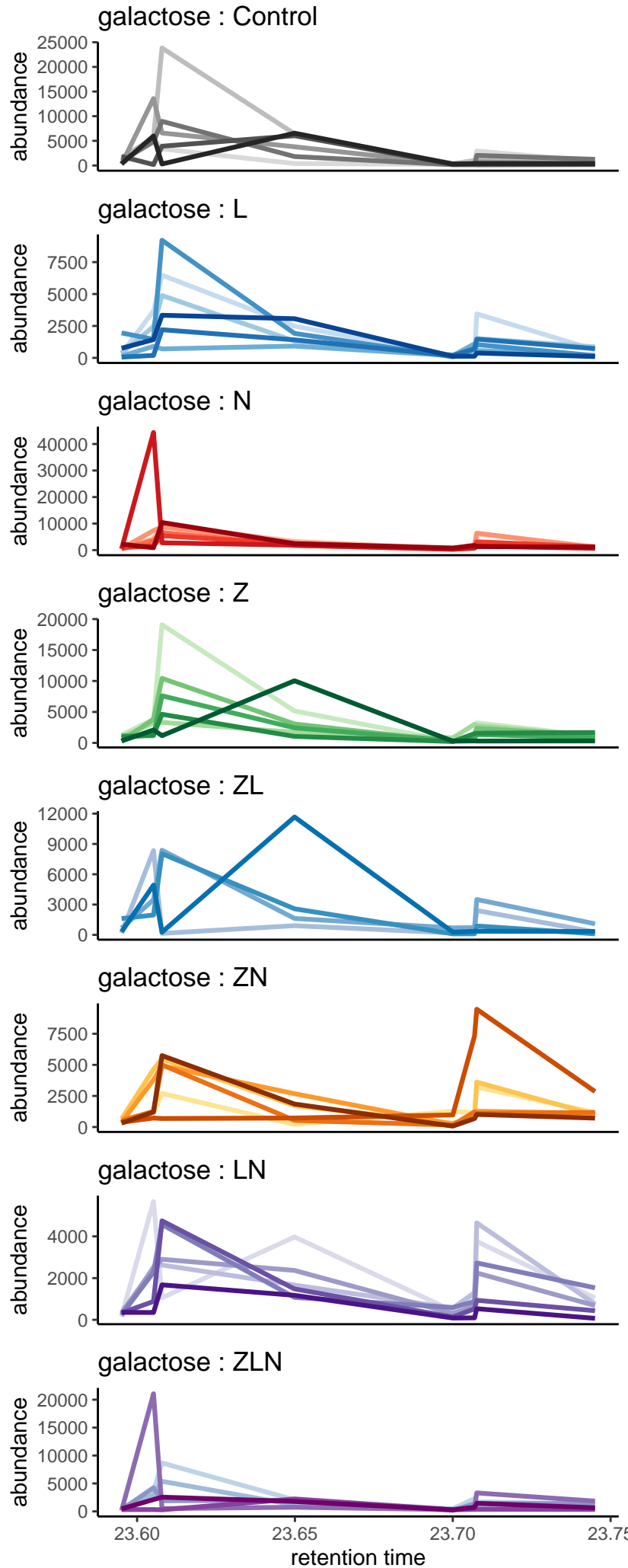


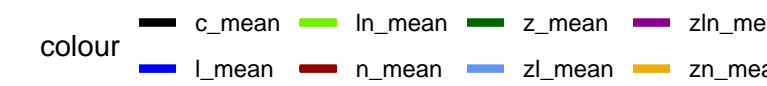
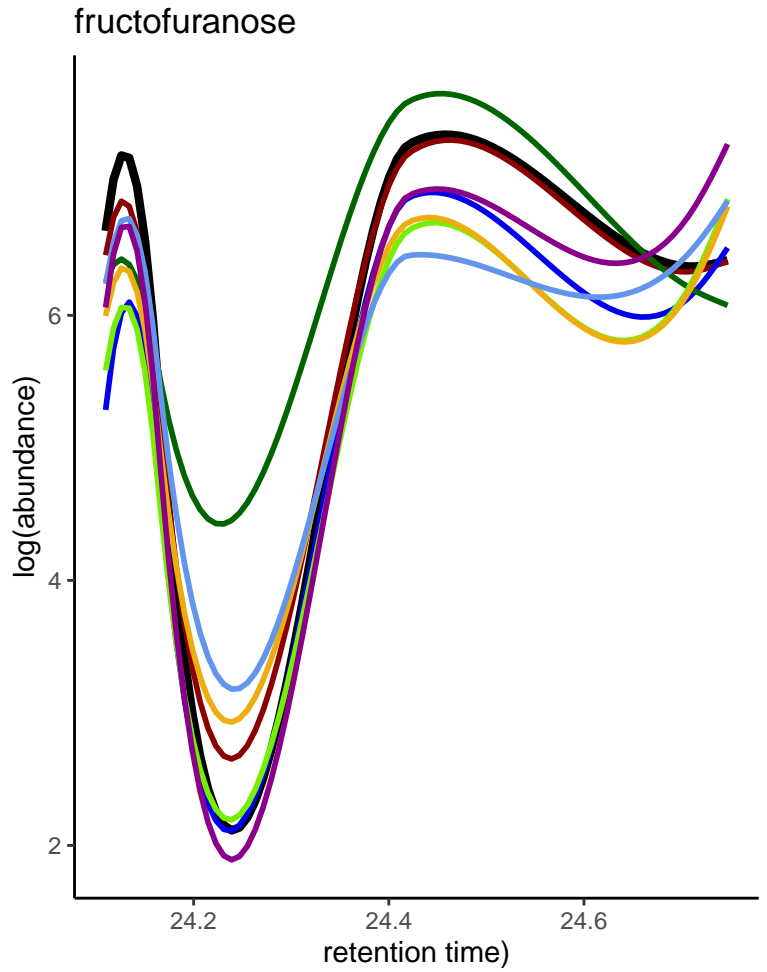
myo-inositol								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
1754	23.21418	0.64	0.03	0.12	0.19	0.06	0.21	1.00
1756	23.21487	0.83	0.01	0.06	0.55	0.09	0.24	0.91
1757	23.21523	0.58	0.01	0.42	0.94	0.06	0.94	0.68
1758	23.21562	0.69	0.01	0.28	0.14	0.09	0.40	0.68
1759	23.22083	0.33	0.03	0.67	0.39	0.00	0.41	0.40
1762	23.22158	0.02	0.07	0.48	0.68	0.07	0.64	0.23
1764	23.22188	0.41	0.52	0.05	0.36	0.02	0.08	0.70
1768	23.22584	0.37	0.01	0.69	0.55	0.29	0.11	0.08
1770	23.23656	0.50	0.01	0.99	0.30	0.06	0.08	0.65
1771	23.24742	0.31	0.01	0.16	0.01	0.40	0.23	0.07



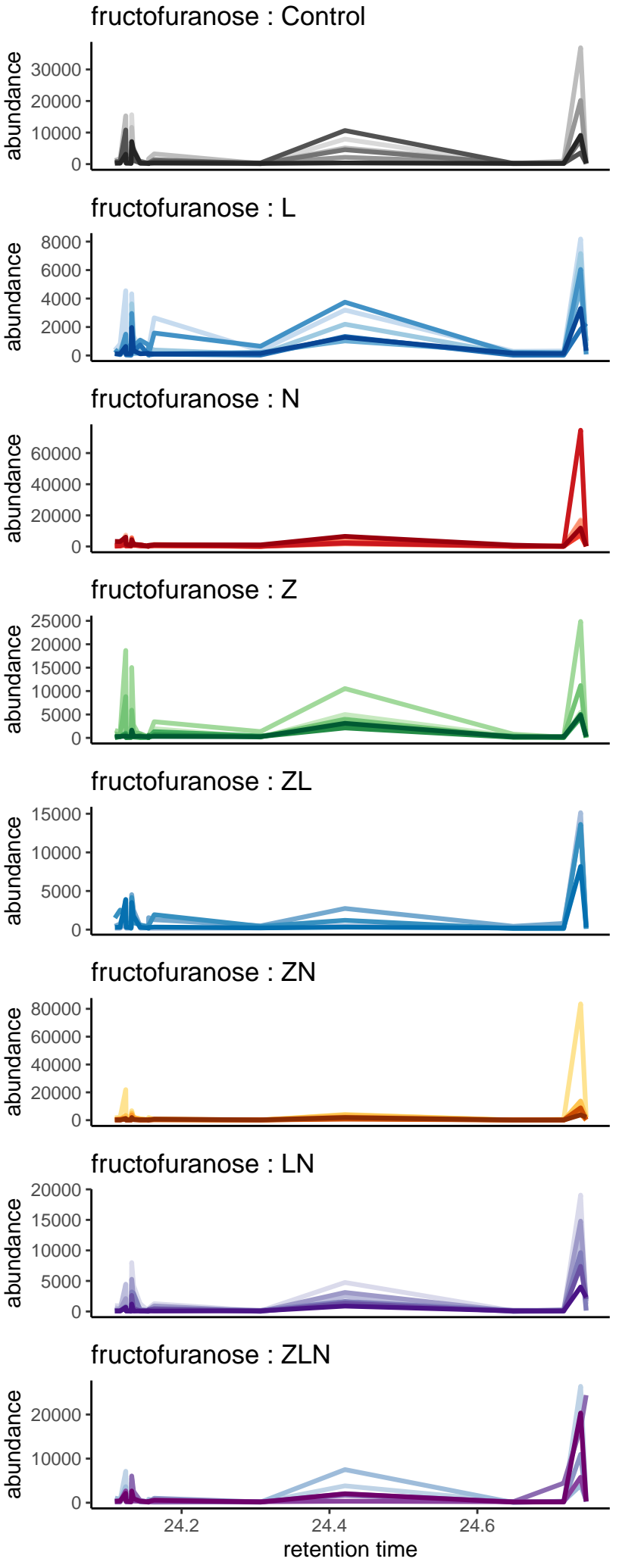


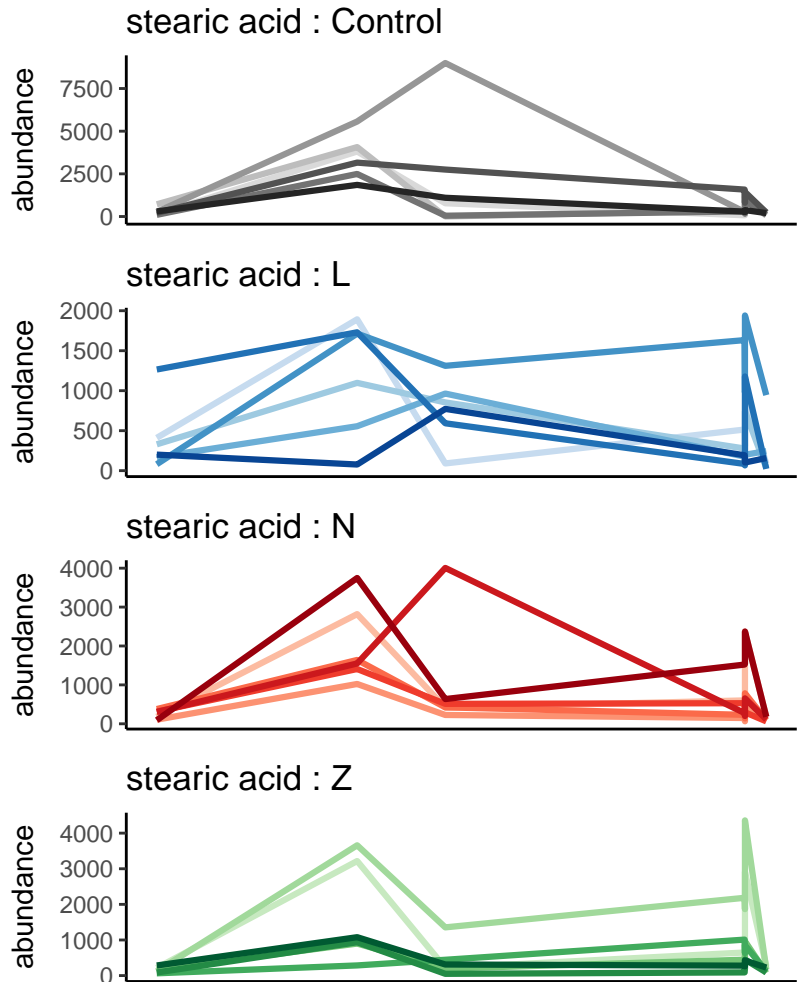
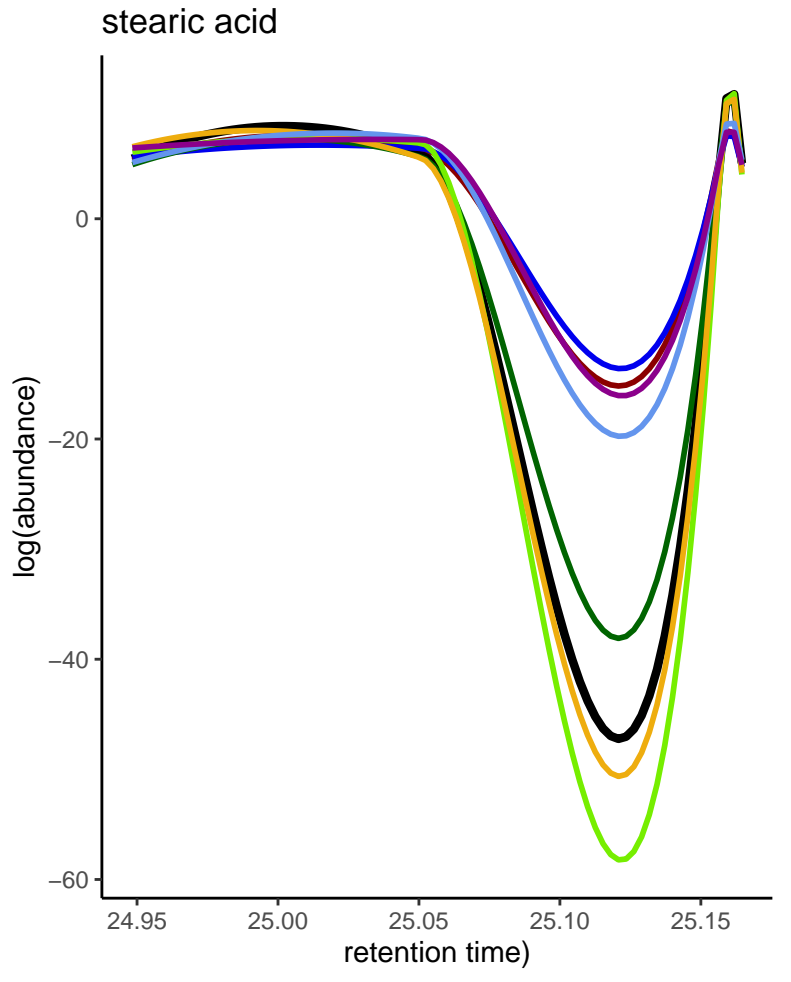
galactose								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
1778	23.59508	0.62	0.02	0.42	0.35	0.44	0.61	0.50
1779	23.60520	0.90	0.39	0.93	0.02	0.36	0.92	0.91
1780	23.60788	0.27	0.02	0.96	0.53	0.75	1.00	0.25
1781	23.64988	0.40	0.48	0.05	0.18	0.09	0.32	0.66
1784	23.70002	0.51	0.11	0.04	0.33	0.19	0.76	0.77
1785	23.70691	0.13	0.04	0.01	0.33	0.63	0.68	0.07
1787	23.70760	0.89	0.37	0.03	0.48	0.18	0.28	0.78
1790	23.74507	0.06	0.02	0.01	0.72	0.93	0.60	0.26





fructofuranose								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zn_fvalue
1800	24.10981	0.96	0.15	0.78	0.05	0.91	0.91	0.80
1801	24.11690	0.42	0.03	0.95	0.02	0.56	0.82	0.58
1811	24.12445	0.94	0.04	0.75	0.03	0.77	0.23	0.94
1814	24.12493	0.86	0.28	0.95	0.05	0.92	0.39	0.63
1828	24.13168	0.74	0.03	0.28	0.01	0.54	0.49	0.14
1833	24.13250	0.29	0.31	0.45	0.03	0.92	0.15	0.26
1841	24.13493	0.79	0.74	0.83	0.02	0.93	0.50	0.02
1846	24.14397	0.36	0.15	0.64	0.02	0.76	0.37	0.25
1851	24.15522	0.21	0.62	0.05	0.03	0.41	0.33	1.00
1852	24.15528	0.07	0.09	0.19	0.02	0.61	0.79	0.67
1854	24.15532	0.96	0.36	1.00	0.03	1.00	0.76	0.19
1856	24.16293	0.76	0.04	0.40	0.62	0.22	0.60	0.94
1858	24.30675	0.08	0.69	0.00	0.22	0.45	0.89	0.26
1860	24.42089	0.32	0.01	0.91	0.82	0.94	0.12	0.13
1870	24.64858	0.28	0.23	0.00	0.02	0.21	0.86	0.07
1871	24.71680	0.55	0.62	0.72	0.00	0.83	0.24	0.90
1877	24.73965	0.81	0.41	0.10	0.05	0.36	0.94	0.41
1879	24.74717	0.52	0.00	0.04	0.13	0.06	0.88	0.58

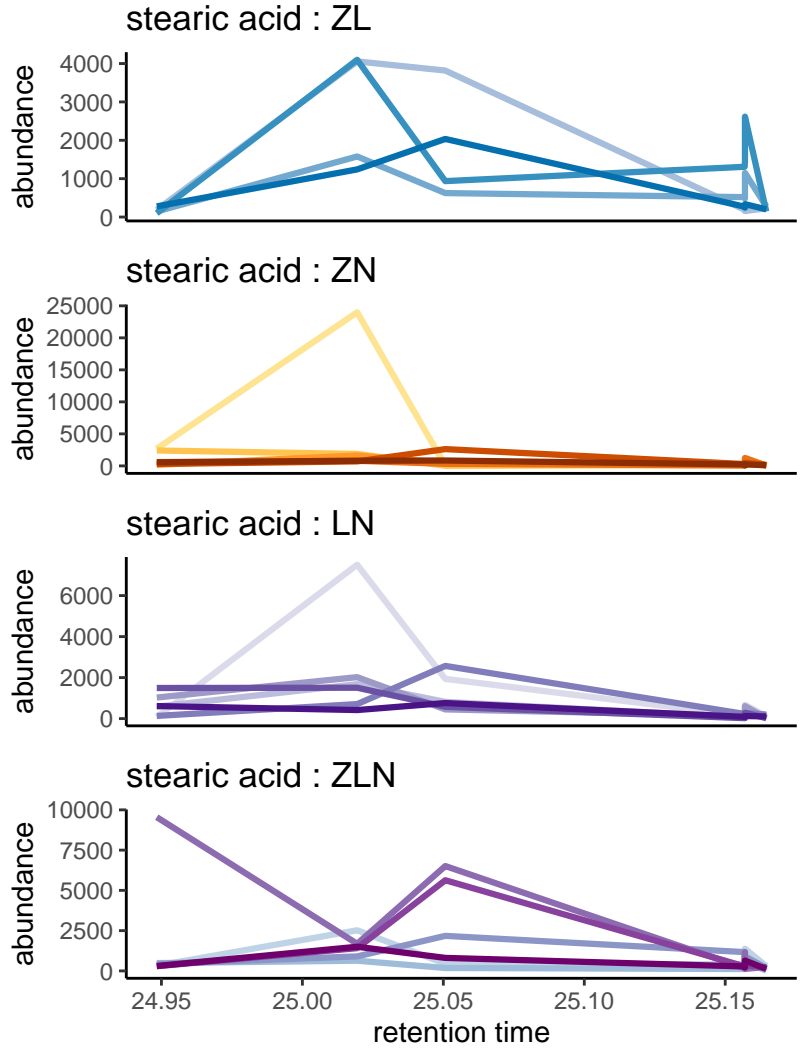


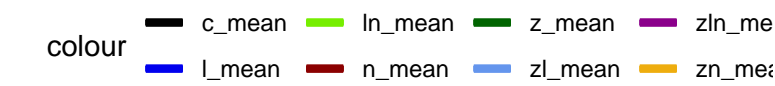
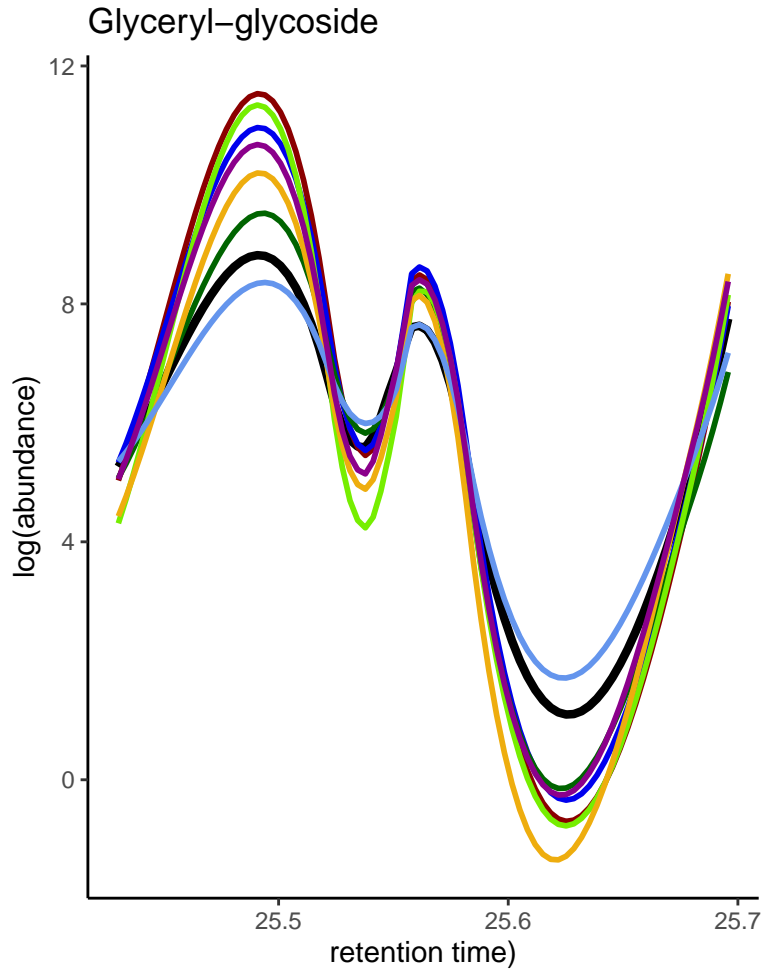


colour

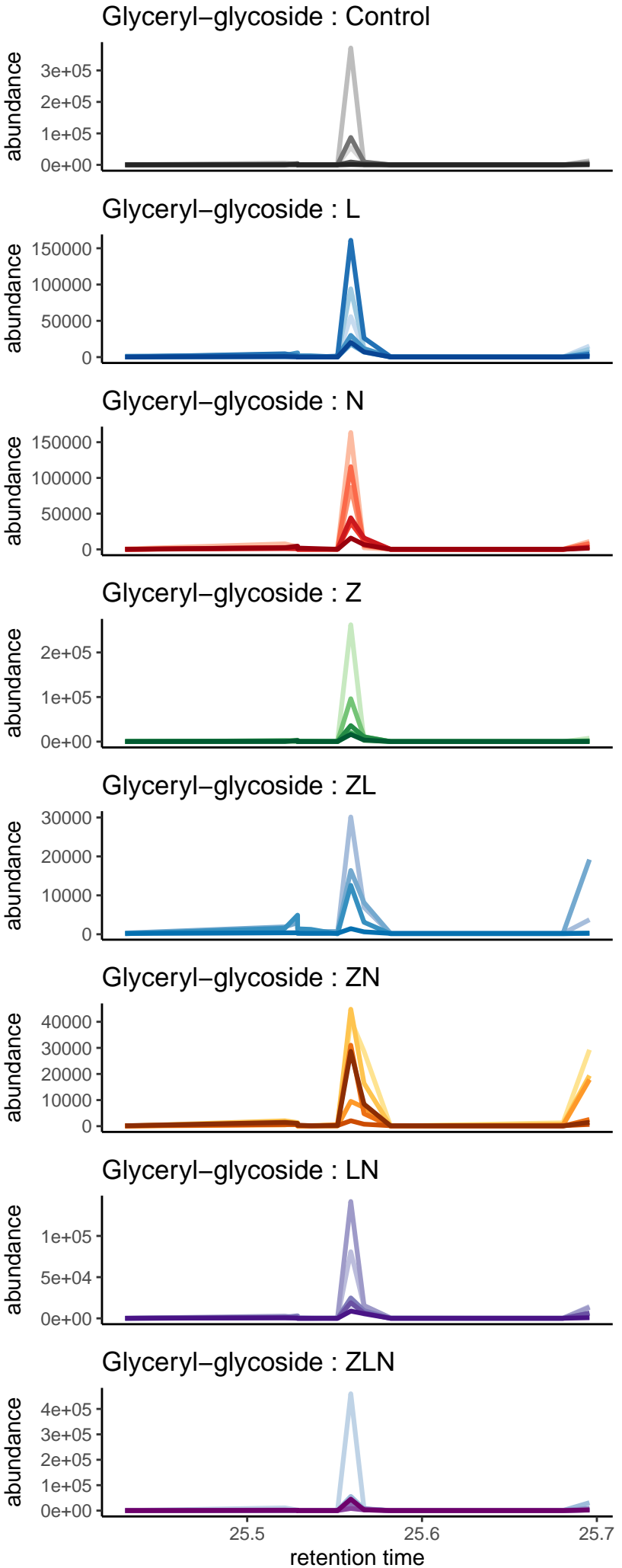
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<span style="color: blue;">—</span> l_mean	<span style="color: darkred;">—</span> n_mean	<span style="color: lightblue;">—</span> zl_mean	<span style="color: orange;">—</span> zn_me

stearic acid								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
1882	24.94834	0.47	0.25	0.01	0.51	0.02	0.77	0.44
1884	25.01943	0.84	0.18	0.96	0.08	0.89	0.74	0.05
1888	25.05070	0.84	0.03	0.61	0.20	0.54	1.00	0.98
1890	25.15688	0.37	0.26	0.03	0.05	0.53	0.48	0.04
1891	25.15700	0.11	0.37	0.00	0.01	0.30	0.52	0.03
1892	25.15700	0.82	0.04	0.14	0.41	0.73	0.93	0.38
1893	25.16457	0.47	0.94	0.03	0.04	0.84	0.95	0.47

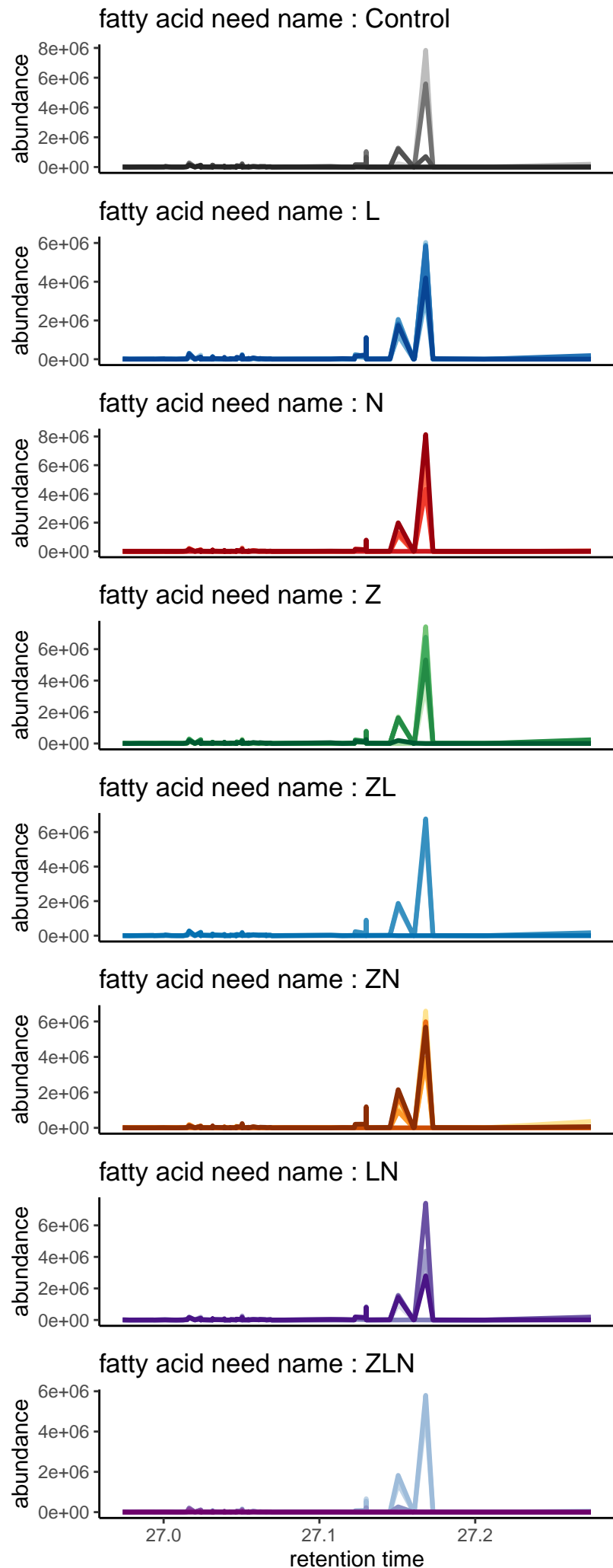
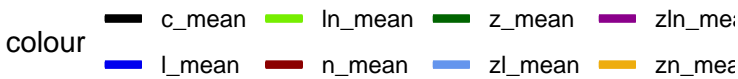




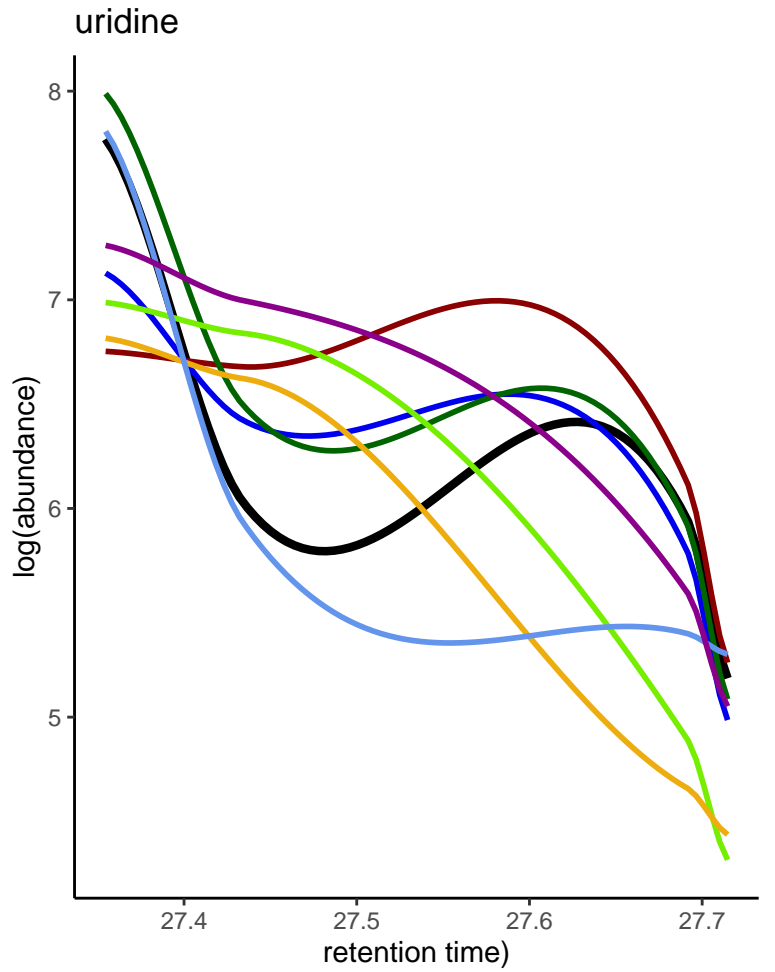
Glyceryl-glycoside								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
1900	25.43023	0.87	0.90	0.03	0.10	0.78	0.59	0.25
1909	25.52154	0.24	0.70	0.33	0.87	0.86	0.33	0.02
1911	25.52883	0.90	0.05	0.04	0.31	0.74	0.47	0.13
1912	25.52883	0.48	0.58	0.07	0.38	0.34	0.05	0.29
1915	25.52890	0.42	0.67	0.01	0.35	0.59	0.24	0.12
1917	25.53644	0.45	0.80	0.01	0.27	0.15	0.12	0.06
1922	25.55167	0.61	0.48	0.34	0.19	0.05	0.71	0.82
1928	25.55926	0.49	0.94	0.44	0.72	0.81	0.78	0.02
1939	25.56683	0.99	0.18	0.15	0.02	0.51	0.18	0.12
1966	25.58203	0.15	0.32	0.07	0.04	0.76	0.58	0.59
1972	25.68065	0.03	0.79	0.56	0.63	0.09	0.62	0.24
1973	25.69578	0.32	0.53	0.03	0.94	0.11	0.78	0.84







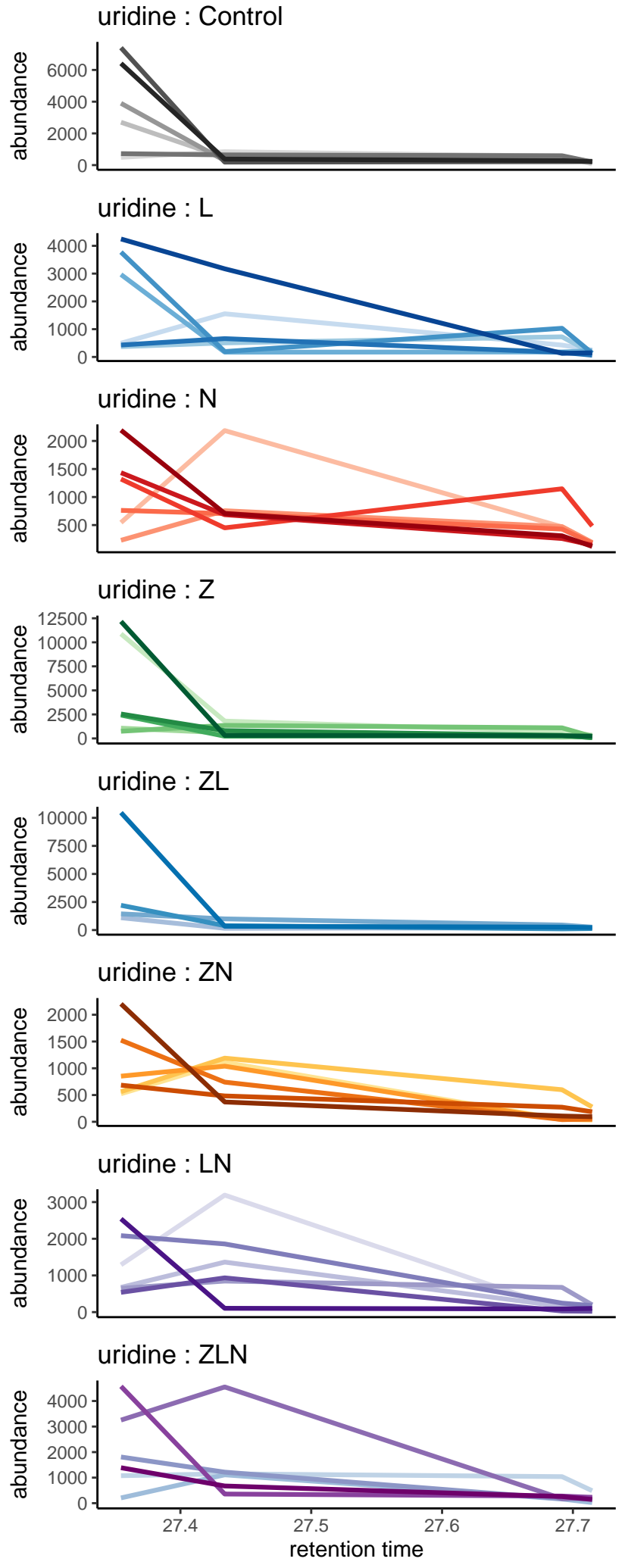


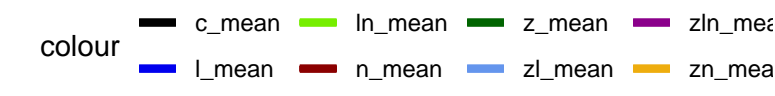
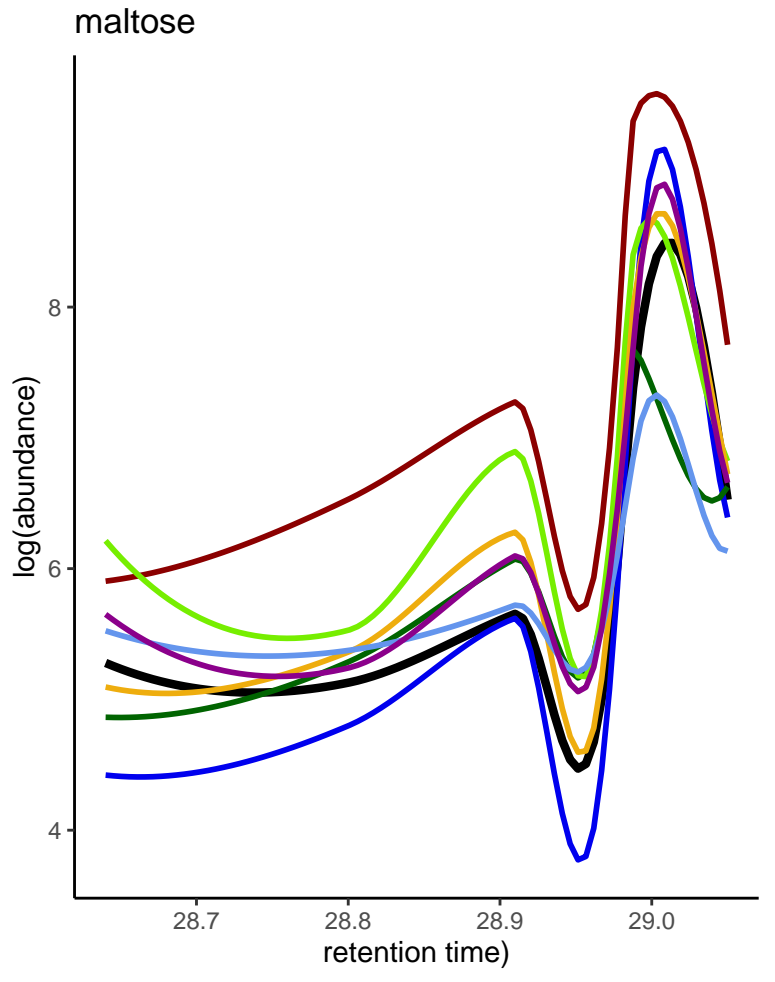


colour

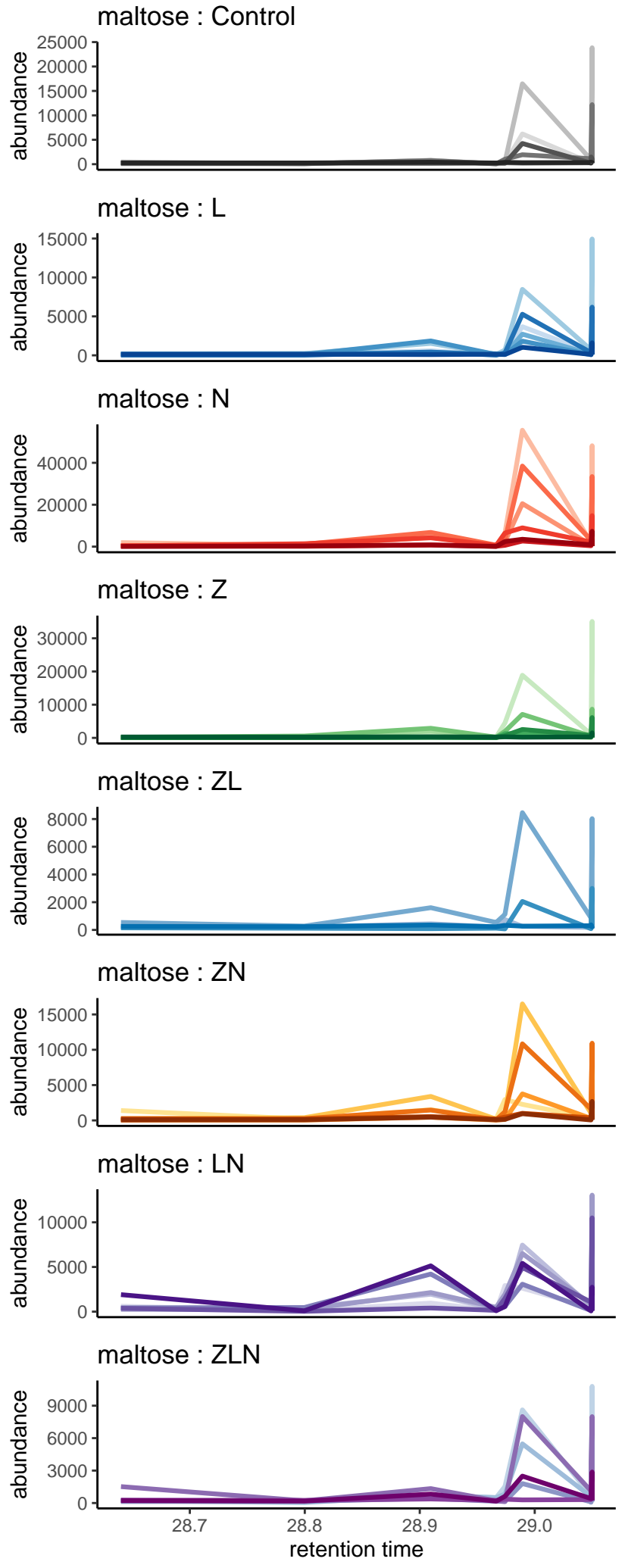
c\_mean ln\_mean z\_mean zln\_mean  
l\_mean n\_mean zl\_mean zn\_mean

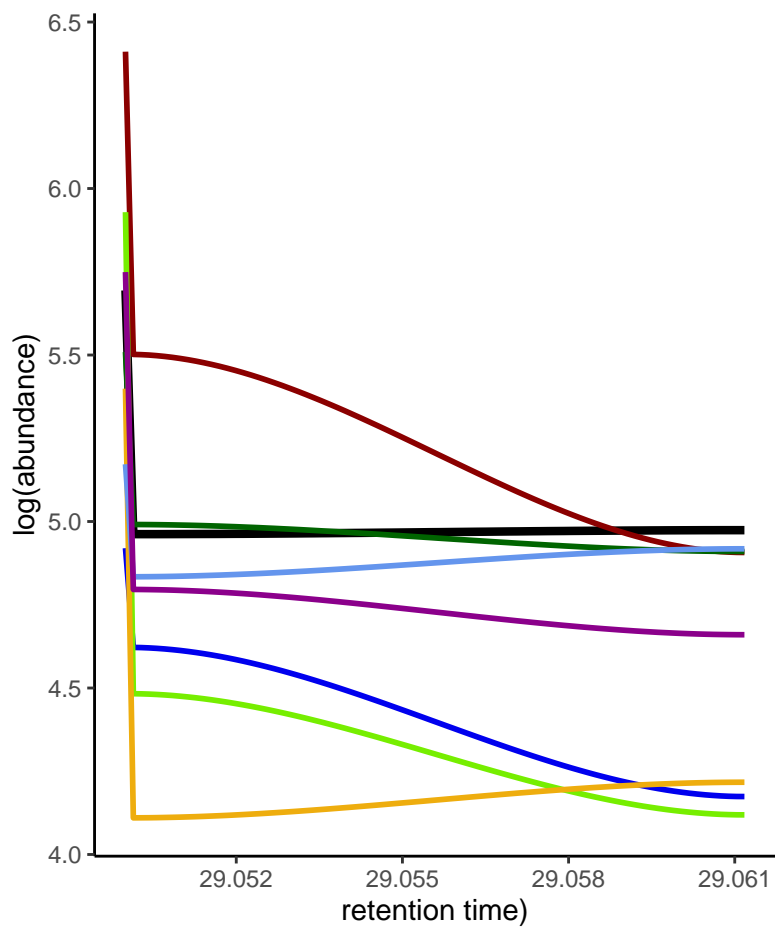
uridine								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
2416	27.35459	0.34	0.81	0.02	0.54	0.59	0.20	0.83
2420	27.43388	0.77	0.54	0.04	0.52	0.98	0.52	0.24
2437	27.69168	0.23	0.30	0.06	0.05	0.60	0.81	0.01
2440	27.71473	0.99	0.59	0.10	0.02	0.68	0.63	0.18





maltose								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
2446	28.64014	0.43	0.54	0.00	0.13	0.05	0.27	0.22
2447	28.79958	0.64	0.07	0.07	0.07	0.02	0.24	0.51
2448	28.90961	0.11	0.37	0.00	0.51	0.07	0.69	0.52
2449	28.96645	0.34	0.90	0.04	0.01	0.02	0.69	0.65
2450	28.97412	0.43	0.03	0.00	0.99	0.00	1.00	0.29
2451	28.98927	0.10	0.47	0.02	0.84	0.58	0.48	0.22
2453	29.04898	0.46	0.04	0.04	0.15	0.29	0.88	0.28
2455	29.04906	0.10	0.13	0.03	0.78	0.52	0.61	0.58
2458	29.04993	0.18	0.44	0.07	0.46	0.30	0.75	0.04
2460	29.04997	0.65	0.19	0.02	0.15	0.49	0.55	1.00

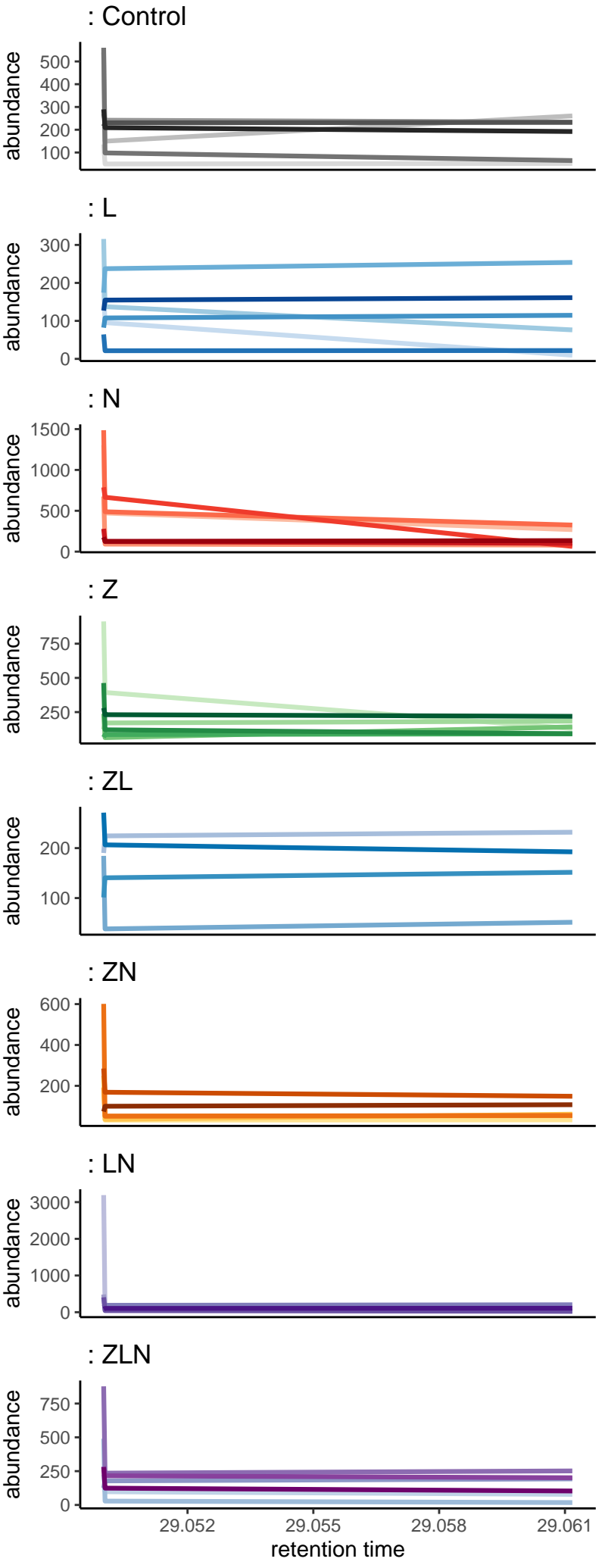


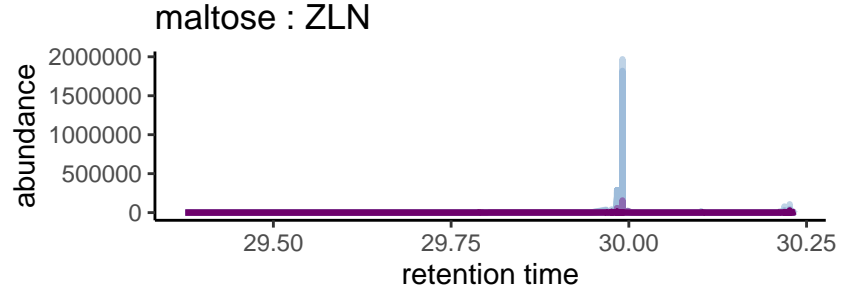
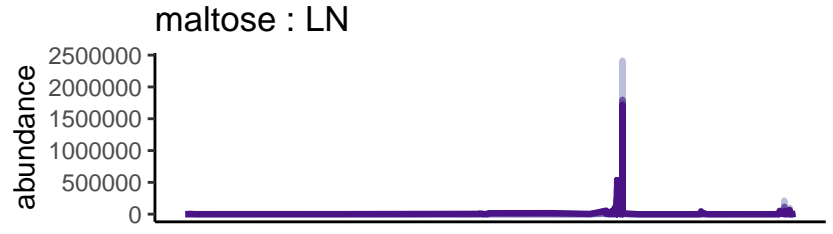
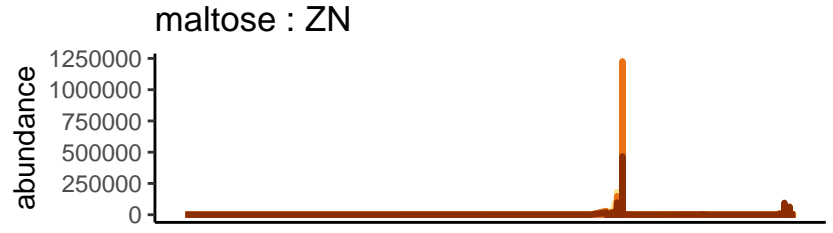
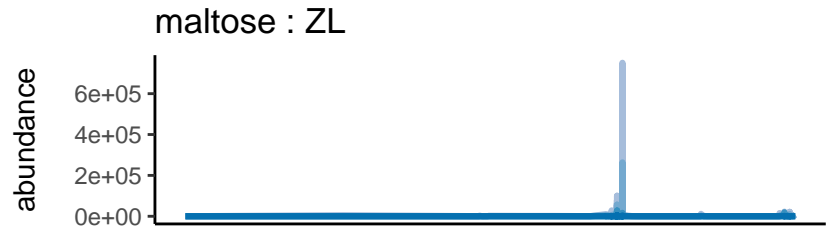
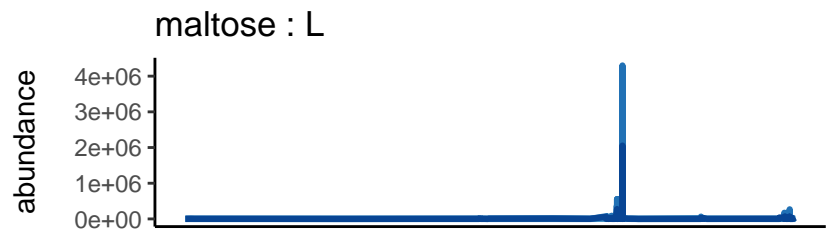
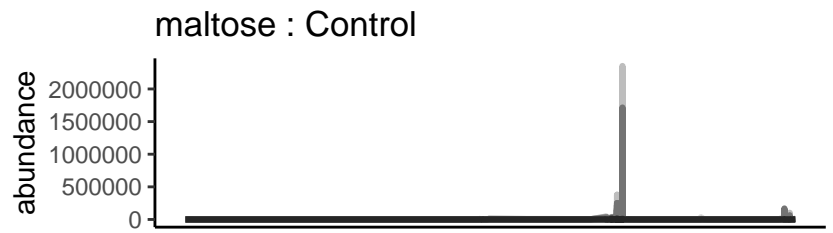
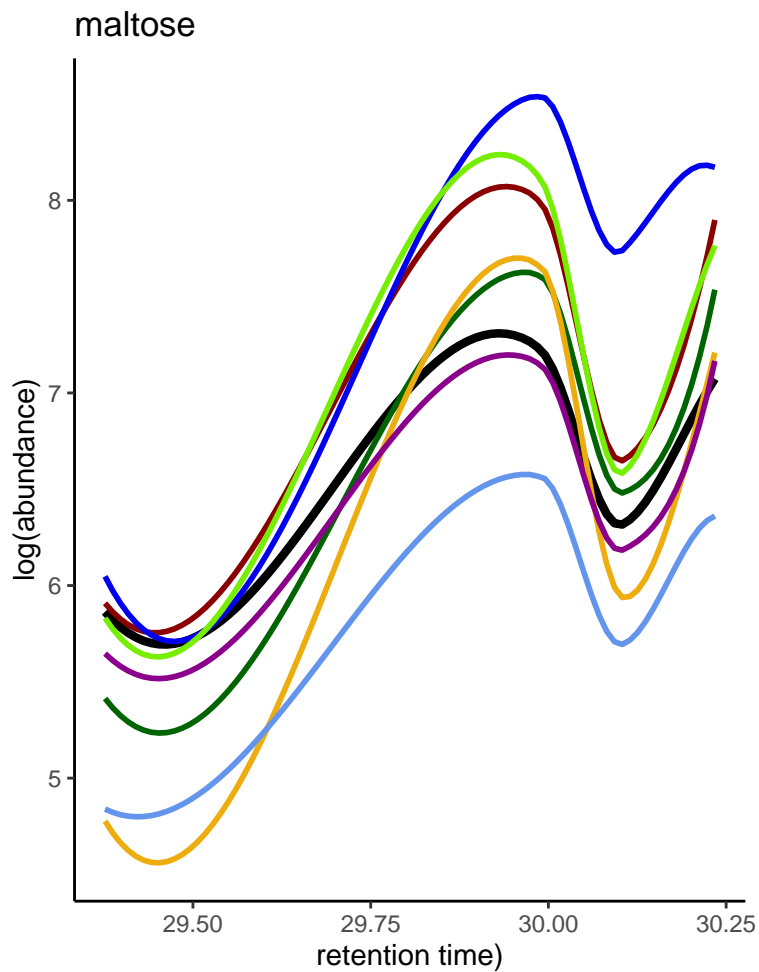


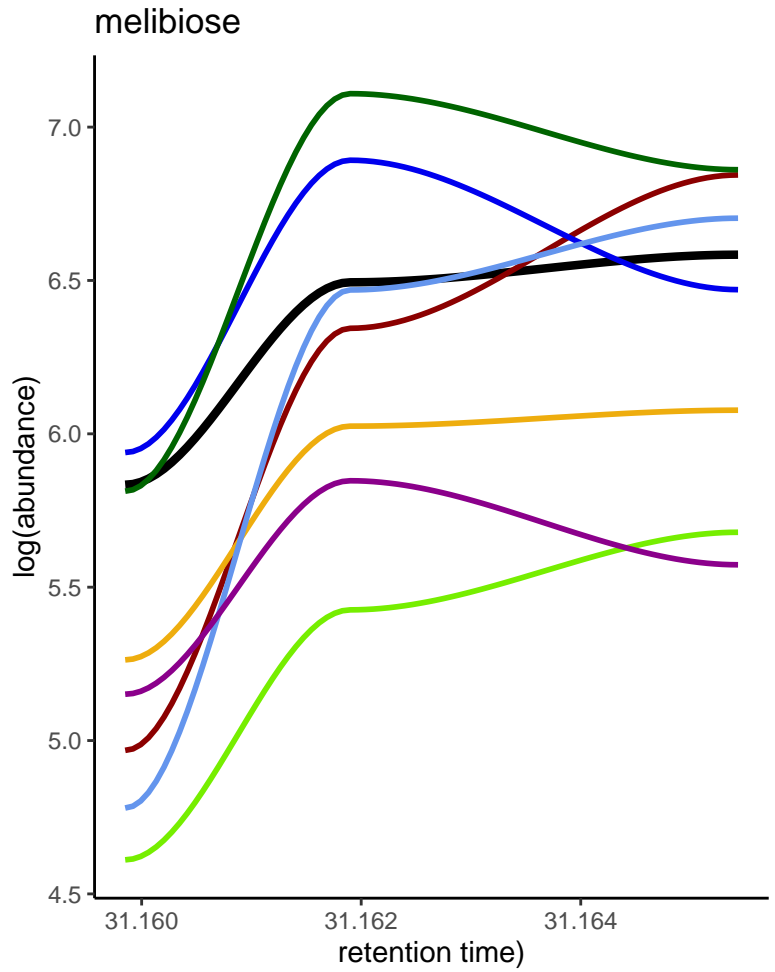
colour

c\_mean ln\_mean z\_mean zln\_mean  
l\_mean n\_mean zl\_mean zn\_mean

X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
2461	29.05000	0.28	0.21	0.02	0.18	0.17	0.32	0.68
2462	29.05003	0.30	0.28	0.61	0.03	0.11	0.93	0.08
2464	29.06117	0.65	0.19	0.32	0.04	0.37	0.67	0.66



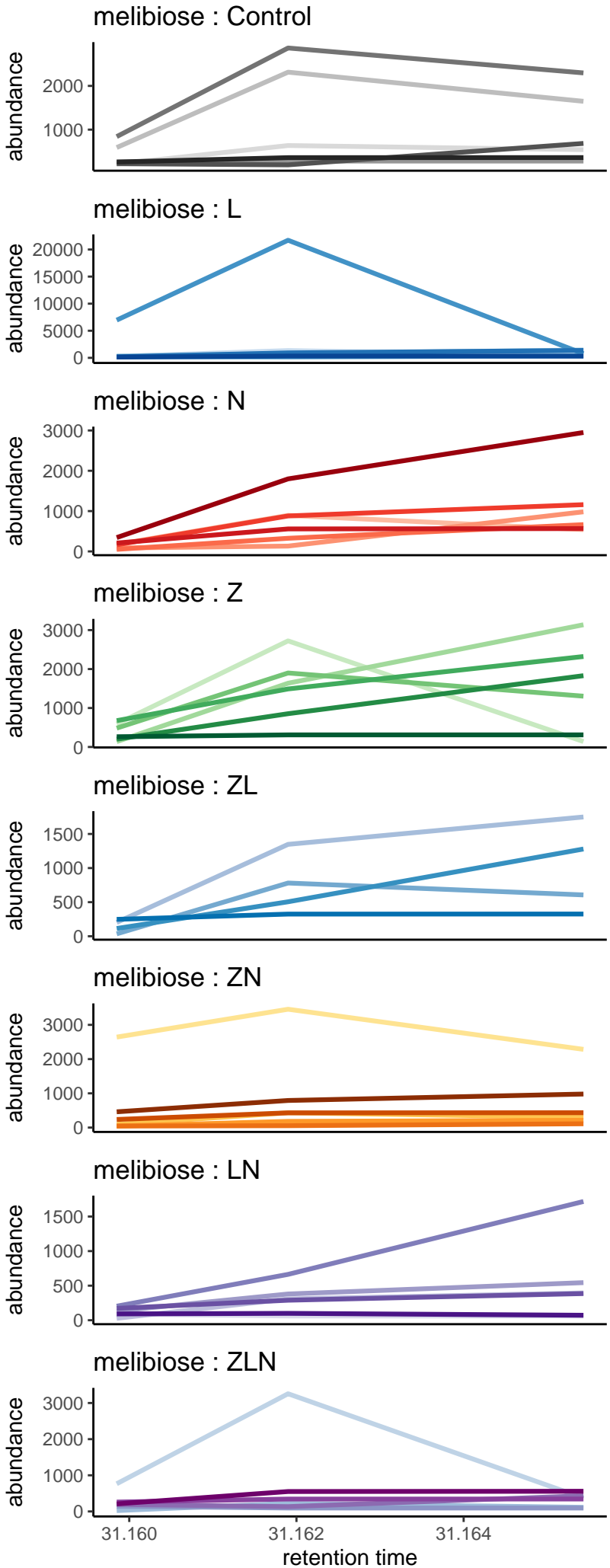


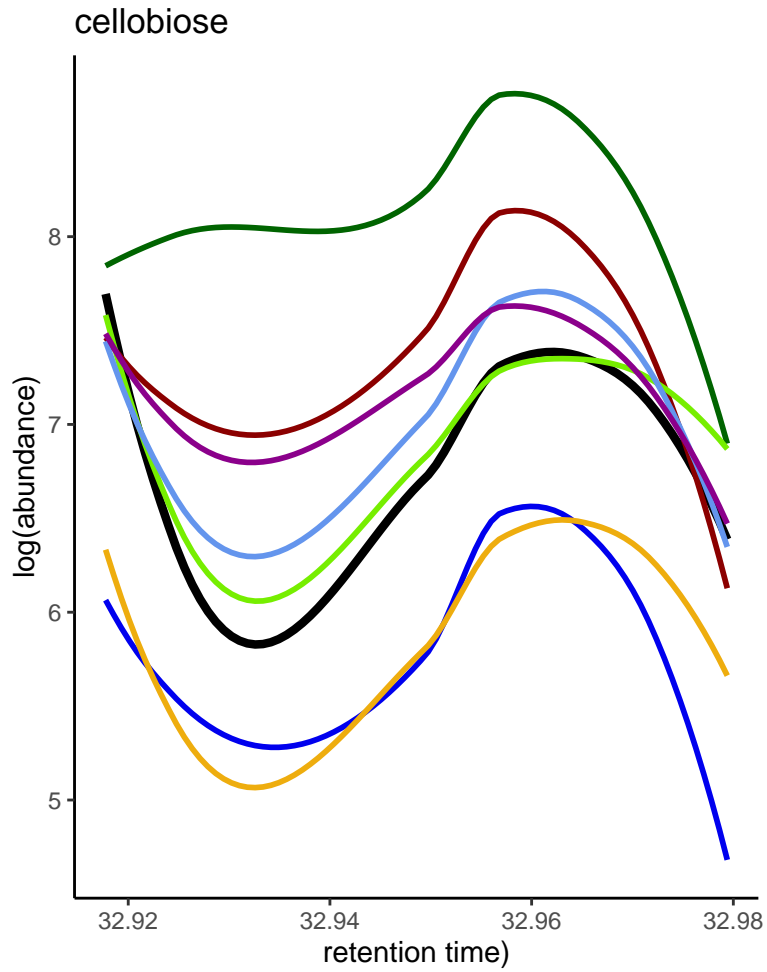


colour

- c\_mean
- ln\_mean
- z\_mean
- zln\_me
- l\_mean
- n\_mean
- zl\_mean
- zn\_me

melibiose								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
2906	31.15985	0.89	0.30	0.04	0.58	0.13	0.76	0.27
2907	31.16190	0.85	0.30	0.02	0.85	0.92	0.49	0.21
2908	31.16543	0.65	0.06	0.05	0.65	0.24	0.21	0.53

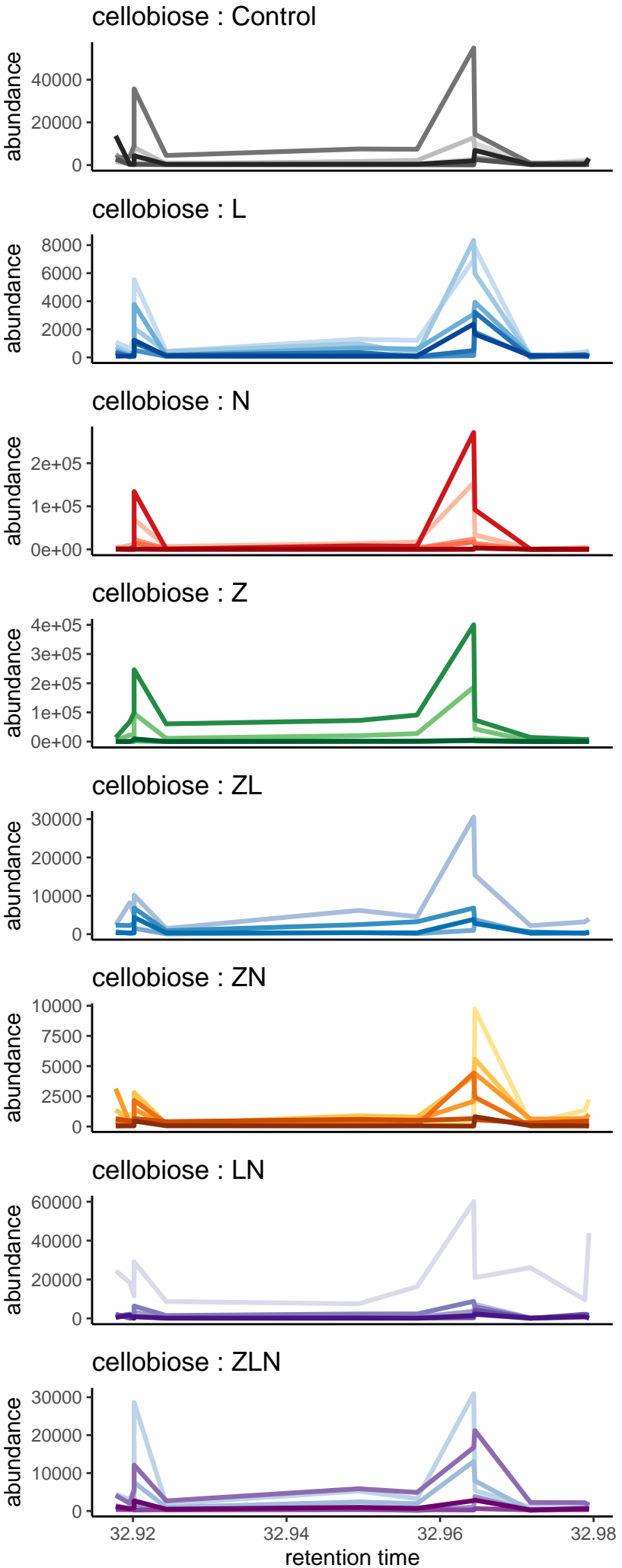


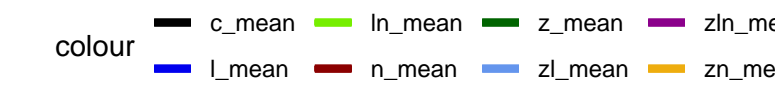
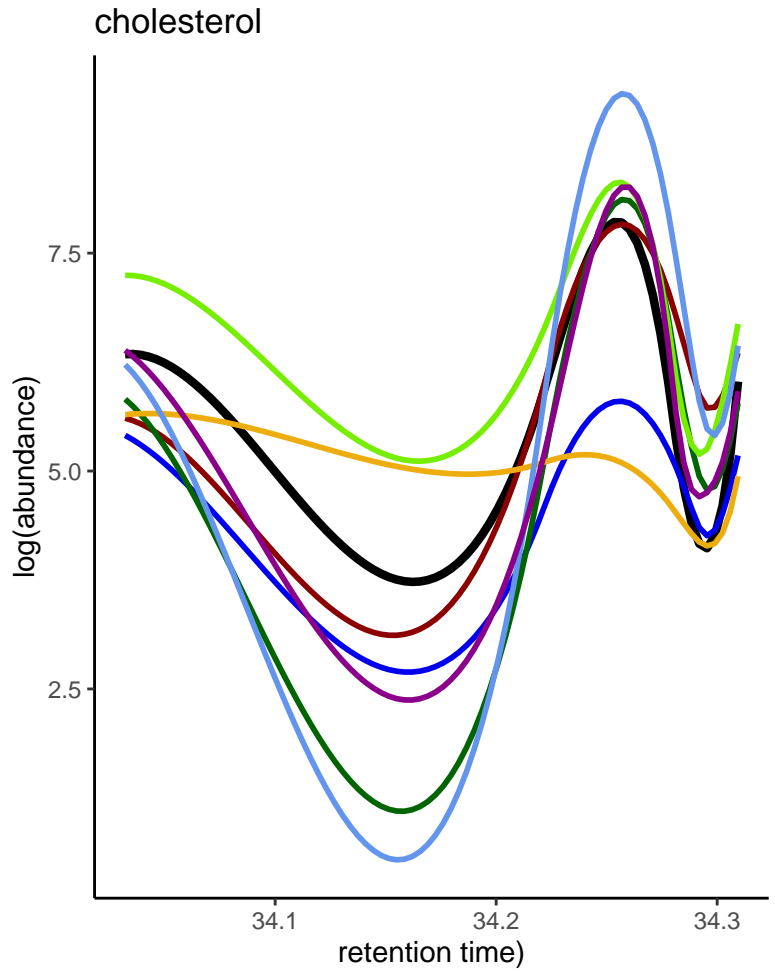


colour

c_mean	ln_mean	z_mean	zln_mean
l_mean	n_mean	zl_mean	zn_mean

cellobiose								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
2917	32.91776	0.39	0.62	0.90	0.07	0.56	0.01	0.22
2919	32.91958	0.50	0.91	0.90	0.31	0.05	0.05	0.82
2920	32.92015	0.16	0.41	0.63	0.36	0.02	0.12	0.16
2921	32.92015	0.78	0.43	0.74	0.38	0.01	0.46	0.10
2922	32.92433	0.64	0.50	0.81	0.26	0.07	0.03	0.32
2925	32.94942	0.34	0.49	0.87	0.31	0.03	0.17	0.19
2929	32.95700	0.40	0.41	0.80	0.35	0.04	0.14	0.42
2933	32.96438	0.80	0.66	0.66	0.50	0.03	0.42	0.21
2934	32.96453	0.46	0.14	0.74	0.34	0.04	0.44	0.19
2935	32.97177	0.03	0.35	0.58	0.27	0.09	0.10	0.89
2936	32.97887	0.73	0.72	0.30	0.60	0.02	0.01	0.75
2937	32.97940	0.45	0.61	0.70	0.37	0.06	0.03	0.44





cholesterol								
X	retention_time	z_fvalue	l_fvalue	n_fvalue	zl_fvalue	zn_fvalue	ln_fvalue	zln_fvalue
2941	34.03213	0.59	0.10	0.31	0.72	0.30	0.01	0.05
2945	34.22190	0.12	0.95	0.19	0.30	0.04	0.25	0.05
2946	34.27983	0.96	0.68	0.82	0.19	0.03	0.95	0.92
2947	34.28532	0.96	0.84	0.90	0.34	0.03	0.53	0.85
2948	34.29257	0.77	0.85	0.59	0.54	0.03	0.35	0.88
2949	34.29322	0.99	0.54	0.36	0.24	0.02	0.92	0.64
2951	34.30823	0.30	0.48	0.67	0.50	0.02	0.55	0.97
2952	34.30946	0.44	0.61	0.47	0.03	0.07	0.14	0.20

