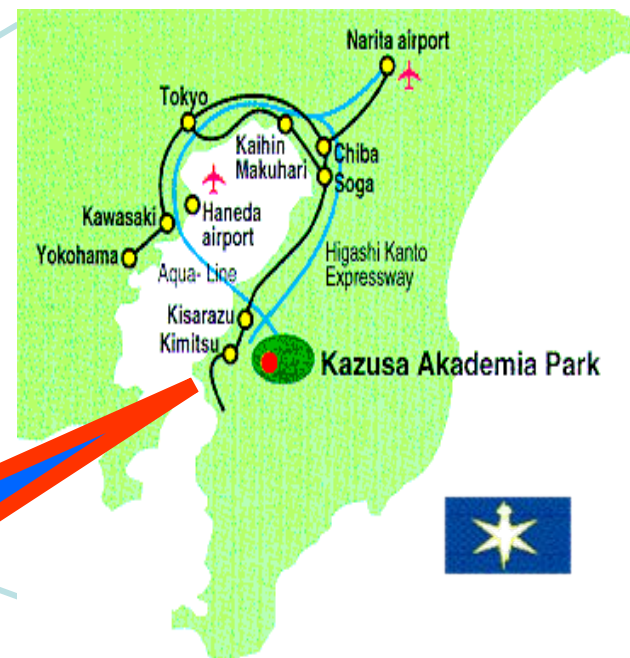


Workshop Argentina-Japan
“Bioscience and Biotechnology for the promotion of
Agriculture and Food Production – August 3rd to 7th 2009-

Metabolomics approaches for Agro-Biotechnology



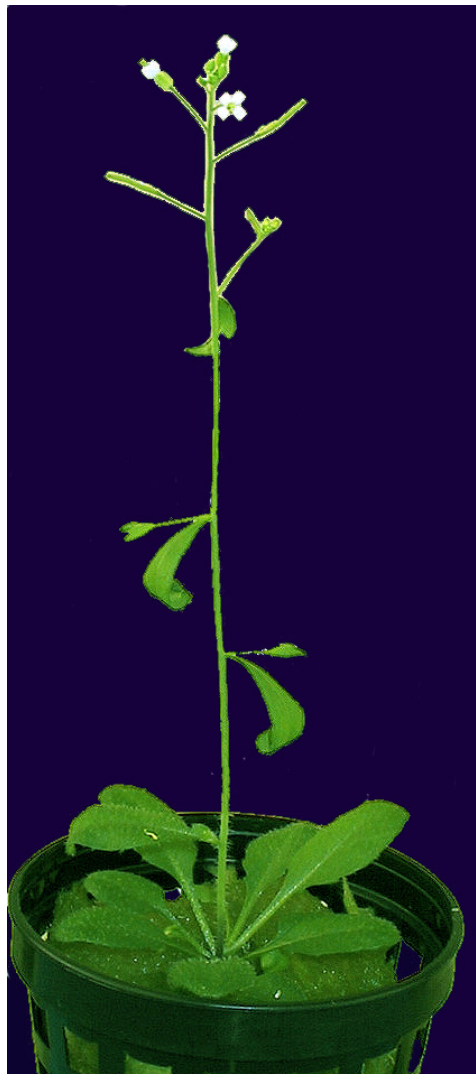
Kazusa DNA Res Inst.
Daisuke Shibata



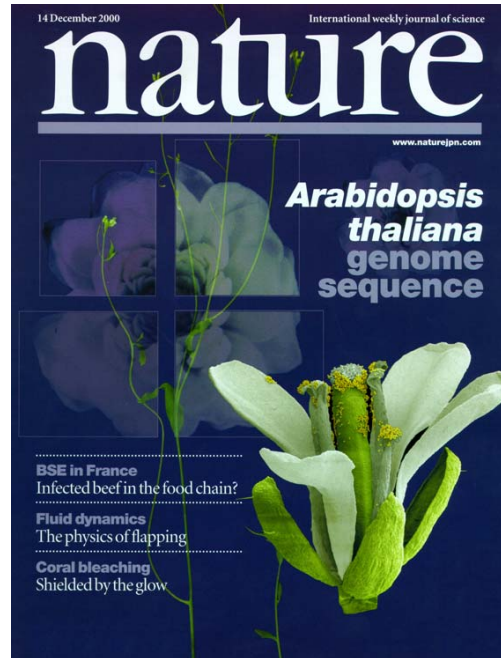
Chiba Prefecture State Government supports financially KDRI.

**Dep. Plant Genomics
Dep. Human Genomics
Dep. Biotechnology**

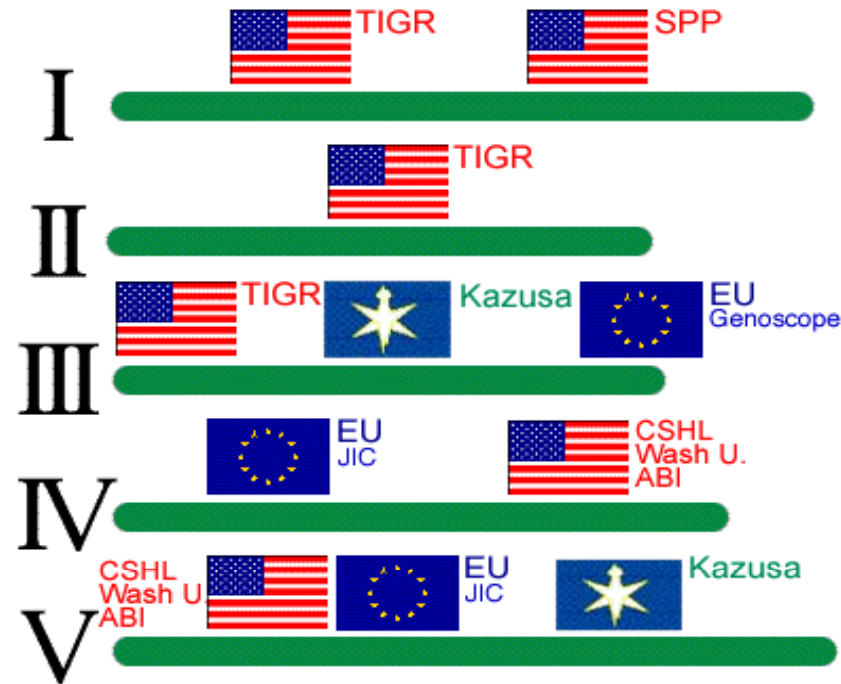
Kazusa DNA Research Institute (KDRI)



Arabidopsis thaliana



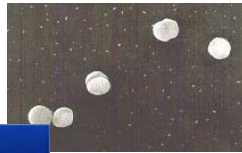
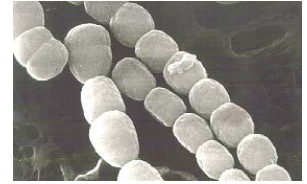
December 2000



Kazusa DNA Res. Inst. (Tabata's lab) sequenced about 30Mb (28% contribution) in the Arabidopsis genome sequencing of 116 Mb.

Plant-related Genome Sequencing at KDRI

Photosynthetic Bacteria	<i>Synechocystis</i> sp. strain PCC 6803	(1996)
	<i>Anabaena</i> sp. PCC 7120	(2001)
	<i>Thermosynechococcus elongatus</i> BP-1	(2002)
	<i>Gloeobacter violaceus</i> PCC7421	(2003)
	→ Photosynthesis	



Nitrogen- fixation bacteria	<i>Mesorhizobium loti</i> MAFF303099	(2000)
	<i>Bradyrhizobium japonicum</i> USDA110	(2002)
	<i>Lotus japonicus</i> (Legume)	(2008)
	→ Nitrogen fixation	



Plants	<i>Arabidopsis</i> (Chromosome 3 &5)	(2000)
	→ Model plant	
	Eucalyptus (for pulp)	In progress
	Tomato (Chromosome 8)	(2009)



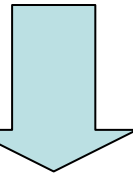
→ Agricultural and industrial application

Metabolomics approaches in agricultural biotechnology

Researches in the Japan Society for Bioscience, Biotechnology, and Agrochemistry are diverged ;

Cop and vegetable breeding
Genetically modified crops
Wood utilization

Fermentation
Food production
More,,,,



“Metabolomics” approaches



High-throughput tools for metabolite analysis



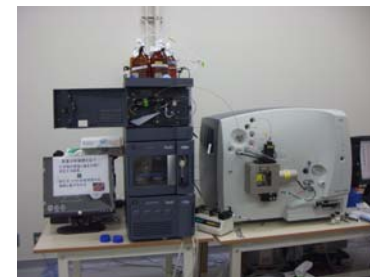
LC-IT-MS



UPLC-Q-TOF-MS



LC-MS



UPLC-TOF-MS



GC-TOF-MS x 2



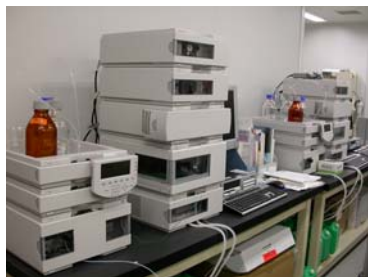
CE-MS x2



FT-IR



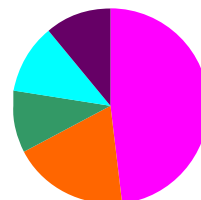
LC-FT/ICR-MS



LC-PDA x2



GPC-MALS

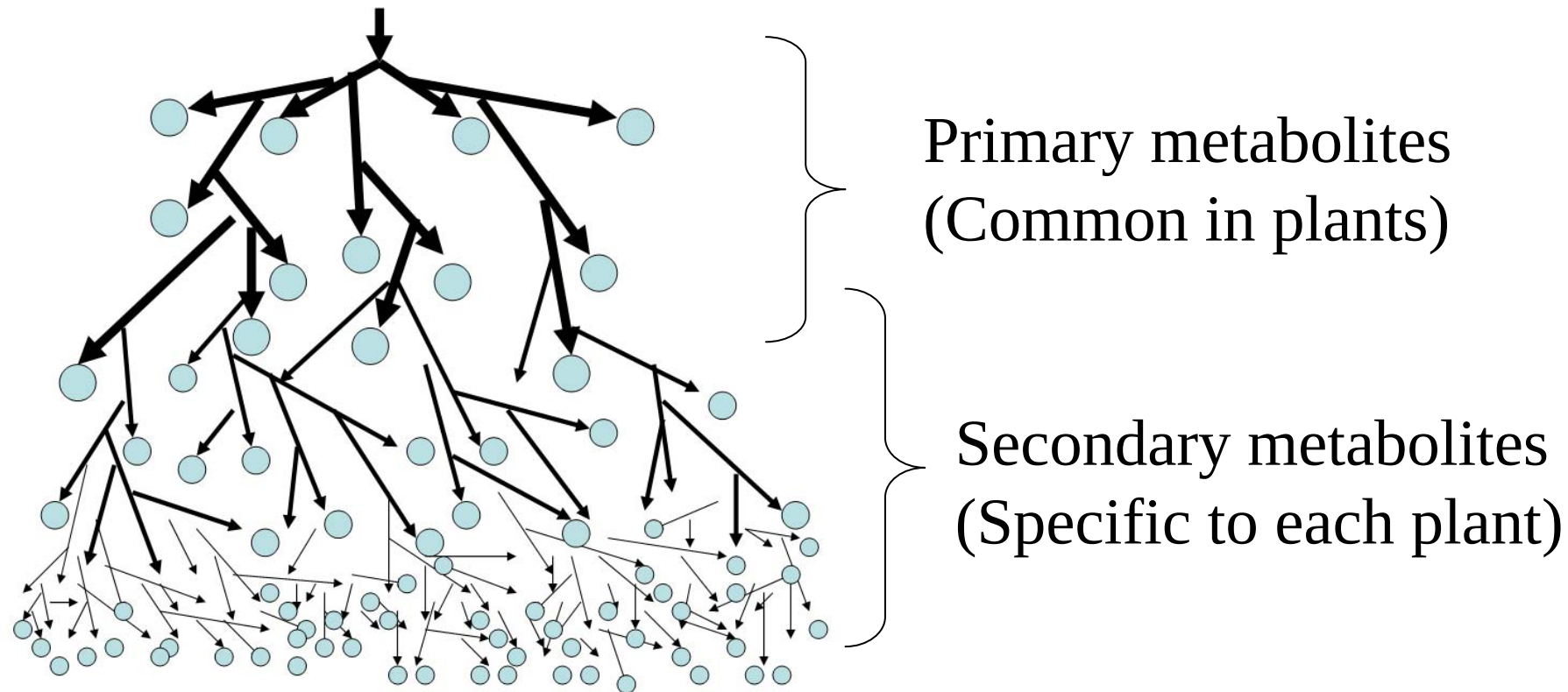


- CE-MS
- GC-TOF-MS
- LC-IT-MS
- LC-FT-MS
- UPLC-MS

of analyzed: > 80,000 at KDRI

What is the problem?

Carbon oxide, water & nutrients



>200,000 metabolites in the plant kingdom

Is it possible to identify all?

Liquid Chromatography Fourier Transform Ion Cyclotron Resonance Mass Spectrometer

LC-FT/ICR-MS



LTQ-FT
(Thermo Electron Co.)
Magnetic field: 7 T



LC-FT-Orbitrap-MS

- **Ultra-high mass accuracy, ~ 1 ppm**
Using internal standards 0.5~0.2 ppm
- **Coupling LC prior to MS**
 - Prevents ion suppression
 - allows UV/Vis spectrum acquisition
 - Separates isomers
 - allows acquisition of Retention Time
 - RT information is useful
to discriminate
 - isotopic ions
 - adduct ions
 - spontaneously fragmented ions
- **Automated MS/MS acquisition**

Ultra-high mass accuracy measurement



For example:

1ppm accuracy: $m/z=500.0000 \pm 0.0005$

$$^{12}\text{C} = 12$$

$$^{13}\text{C} = 13.003355$$

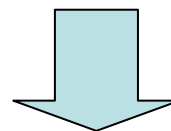
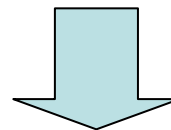
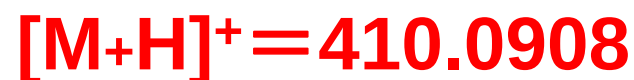
$$^1\text{H} = 1.0078250319$$

$$^{16}\text{O} = 15.9949146223$$

$$^{14}\text{N} = 14.0030740074$$

$$^{31}\text{P} = 30.973763$$

$$^{32}\text{S} = 31.972071$$



MS/MS

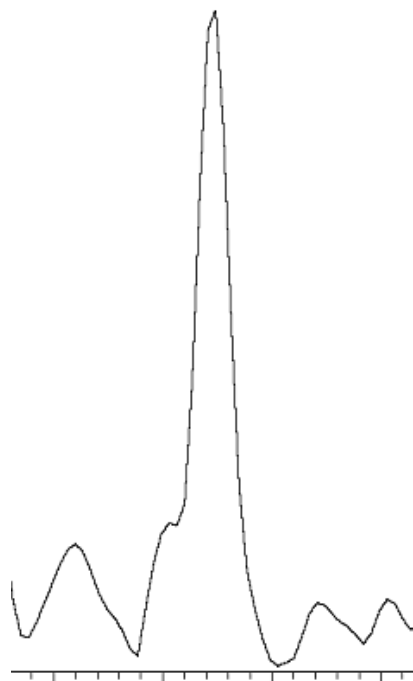
Structure Speculation

Speculation of Molecular Formula by FT-ICR-MS

$^{12}\text{C}=12.000000$ $^1\text{H}=1.007825$, $^{14}\text{N}=14.003074$, $^{16}\text{O}=15.994915$

433.11270

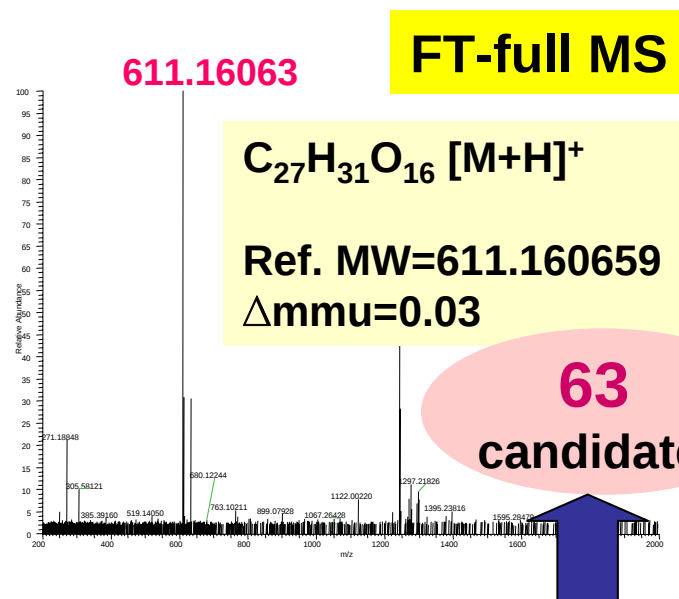
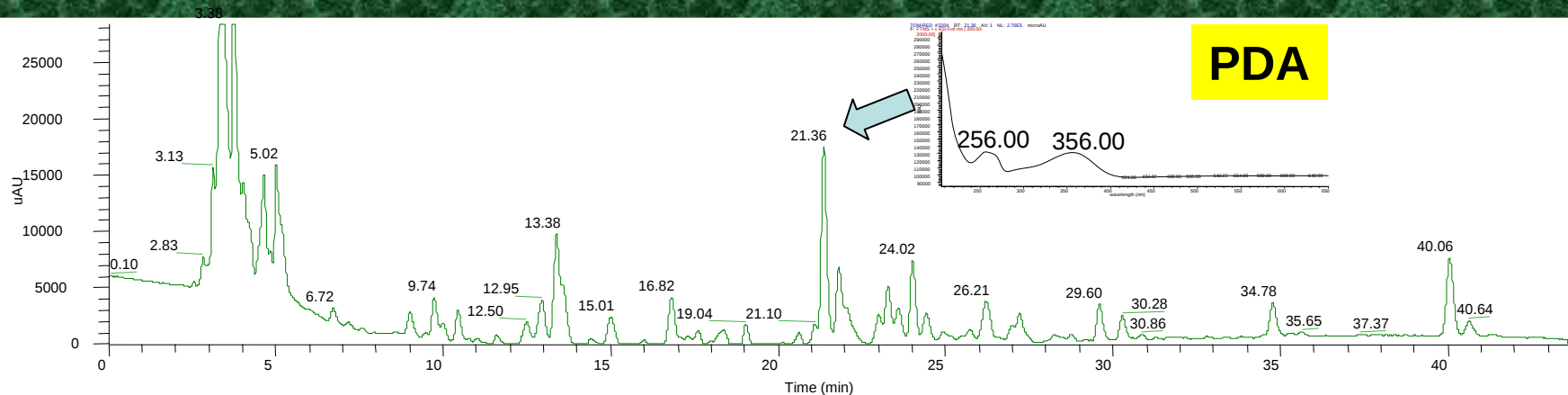
433.11270 $\rightarrow$ $\text{C}_{21}\text{H}_{20}\text{O}_{10}[\text{+H}]^+$



C	H	O	N	S		
21	21	10	0	0	433.112 7	2
13	25	12	2	1	433.112 2	7
11	13	2	16	1	433.112 2	6
14	29	7	2	3	433.113 1	3
11	21	1	12	3	433.111 7	9
•	•	•	•	•	•	
•	•	•	•	•	•	

Milli mass values give a limited number of formulas.

LC-FTICR-MS analysis of tomato fruit



FT-full MS

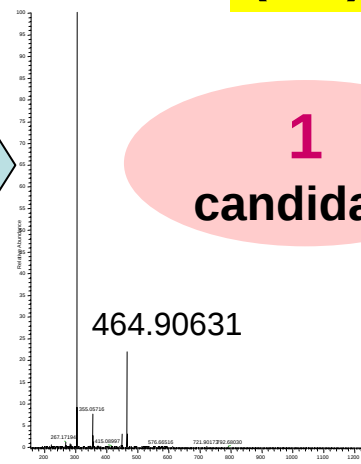
63
candidates

Metabolite
Database

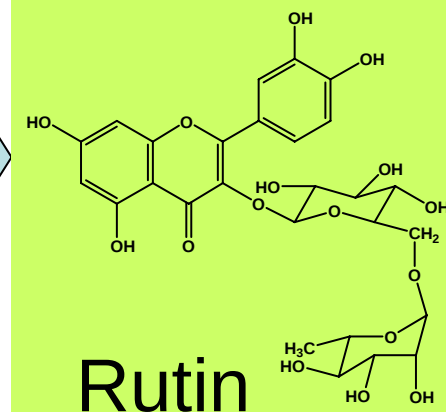
303.09119

(MS)ⁿ

1
candidate

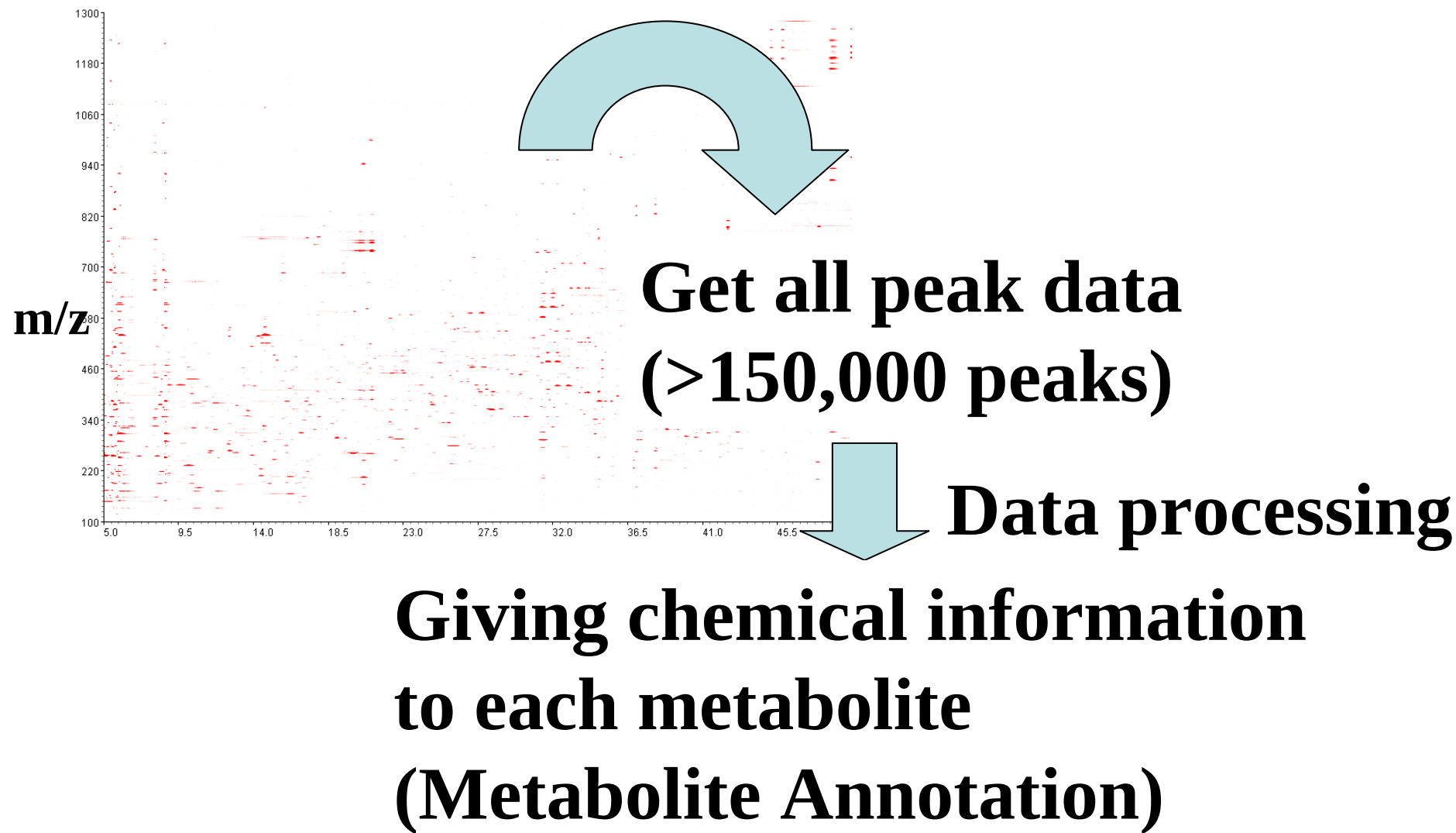


MS/MS Fragmentation
UV absorption (PDA)



Rutin

Annotation, but not identification



Comprehensive analysis of tomato metabolites

Tissues

Fruit 4 stages, 2 tissues



G Y O R

Peel (P), flesh (F)

Leaf, Root, Flower

Extraction

70% methanol

100% chloroform

Ionization

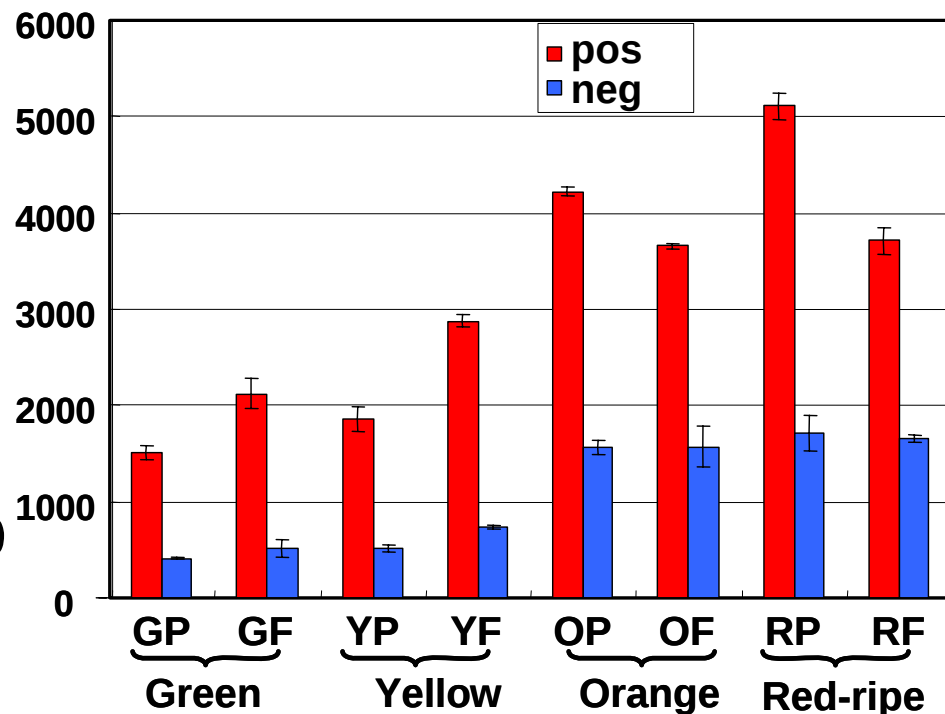
LC C18, C30

ESI (positive, negative)

APPI (positive, negative)

Number of detected peaks

ESI (Methanol fraction)



lijima et al. (2008) Plant J.

How many metabolites ?

70% MeOH
extraction

LC-ESI-based MS

70% MeOH extraction

LC-ESI-based MS

Tissues	Ionization mode	Number of mass signals ^a	Number of peak groups ^a	Number of metabolites ^a	Number of annotated metabolites	Total number of annotated metabolites in each tissue ^b	Annotation grade			
							A	B	C	
	Mature green									
	Flesh	Positive	30 412 ± 3069	1470 ± 155	306 ± 35	154	267			
		Negative	17 292 ± 1483	1673 ± 102	305 ± 22	167				
	Peel	Positive	42 734 ± 5067	2311 ± 260	479 ± 69	228	368			
Negative		20 769 ± 2938	1925 ± 226	397 ± 51	228					
	Breaker									
	Flesh	Positive	28 782 ± 8835	1729 ± 271	357 ± 96	182	291			
		Negative	15 853 ± 4078	1604 ± 311	308 ± 66	168				
	Peel	Positive	43 462 ± 9540	2621 ± 379	636 ± 119	250	440			
Negative		32 675 ± 4440	2733 ± 376	602 ± 85	295					
	Turning									
	Flesh	Positive	24 353 ± 6111	1680 ± 58	352 ± 26	188	284			
		Negative	12 498 ± 4924	1239 ± 460	251 ± 134	156				
	Peel	Positive	63 258 ± 6645	3495 ± 348	784 ± 112	358	611			
Negative		39 274 ± 3449	3187 ± 364	676 ± 79	402					
	Red									
	Flesh	Positive	28 109 ± 1791	1700 ± 132	353 ± 42	179	263			
		Negative	13 808 ± 4403	1444 ± 414	266 ± 64	144				
	Peel	Positive	70 278 ± 3619	4305 ± 288	1039 ± 77	445	696			
Negative		55 429 ± 2452	4723 ± 301	1026 ± 68	428					

Annotation grade

Description of availability of supporting evidence

^aNumbers indicate means ± SD of three measurements.

^bTotal numbers of non-redundant annotated metabolites detected in positive- and negative-ionization modes.

Description of
availability of
supporting evidence

Total: 869 non-redundant metabolites
Including 494 metabolites not in the databases

Annotation of other metabolites Finished at molecular formula-level

Approx. 5600 non-redundant peaks

Checking candidate formulas, providing Evidences.

Detected	Ionization	Formula	GF	GP	YF	YP	OF	OP	RF	RP	Speculated
104.0704	Ep	C4H9NO2									GABA
130.0498	Ep	C5H7NO3									Pyroglutamic acid
134.0449	Ep	C4H7NO4									Aspartic acid
138.0549	Ep	C7H7NO2									p-Aminobenzoic acid
147.0765	Ep	C5H10N2O3									glutamine
147.1128	Ep	C6H14N2O2									L-Lysine
148.0605	Ep	C5H9NO4									L-Glutamic acid
156.0768	Ep	C6H9N3O2									L-Histidine
166.0863	Ep	C9H11NO2									L-Phenylalanine
175.0966	Ep	C8H14O4									Suberic acid
175.1190	Ep	C6H14N4O2									L-arginine
189.1598	Ep	C9H20N2O2									
191.1026	Ep	C7H14N2O4									meso-2,6-Diaminoheptanedioate
193.0344	Ep	C6H8O7									citric acid
205.0971	Ep	C11H12N2O2									Tryptophan
207.1128	Ep	C11H14N2O2									12-Hydroxycytisine
212.0919	Ep	C10H13NO4									
219.1492	Ep	C13H18N2O									
230.0151	Ep	C5H11NO5S2									
245.0769	Ep	C9H12N2O6									5-Ribosyluracil
251.0696	Ep	C8H14N2O5S									
251.1390	Ep	C13H18N2O3									N-Caffeoylputrescine
261.0370	Ep	C6H13O9P									
263.1392	Fn	C14H18N2O3									

Annotation for major secondary metabolite Flavonoids

Rt	Detected	Theoretical	delta ppm	Formula	MS/MS	λ max	GP	GF	YP	YF	OP	OF	RP	RF	
27.06	394.0954	394.0955	-0.21	C18H19O7NS	273(100)	290									
26.56	394.0956	394.0955	-0.54	C18H19O7NS	273(100)	290									
24.02	410.0906	410.0904	0.55	C18H20O8NS	289(100), 375(13)	284									
24.48	410.0907	410.0904	0.62	C18H20O8NS	289(100), 375(13)	284									
33.9	435.1288	435.1286	0.47	C21H22O10	273(100)	284, 336(w)									
34.86	435.1288	435.1286	0.55	C21H22O10	273(100)	362									
37.19	435.1284	435.1286	-0.29	C21H22O10	273(100)	368									
37.65	435.1286	435.1286	0.55	C21H22O10	273(100)	368									
26.16	449.1077	449.1078	-0.31	C21H20O11	287(100)	256, 354									
33.57	449.1083	449.1078	1.03	C21H20O11											
35.81	451.1231	451.1235	-0.86	C21H22O11	289(100)	262, 354									
40.29	451.1236	451.1235	0.18	C21H22O11	289(100)	370									
39.31	451.1236	451.1235	0.65	C21H22O11	289(100)	284									
30.63	465.1030	465.1028	0.52	C21H20O12	303(100)	256, 352									
34.52	465.1393	465.1391	0.44	C22H24O11	303(100)	284, 358									
26.49	495.1860	495.1861	-0.23	C24H30O11	289(100), 451(24), 452(5), 477(4)										
37.78	521.1286	521.1290	-0.63	C24H24O13	273(100), 274(11), 297(5), 315(9), 339(5), 381(8), 485(7), 503(14), 504(3)	290, 366									
41	521.1288	521.1290	-0.32	C24H24O13	273(100), 274(12), 297(3), 315(4), 339(2), 381(3), 485(2)	366									
33.42	537.1237	537.1239	-0.34	C24H24O14	289(100), 290(10), 313(2), 331(14), 355(7), 383(4), 397(9), 501(11), 515(9)	284									
38.2	551.1395	551.1395	-0.06	C25H26O14	303(100), 304(13), 327(5), 345(10), 369(5), 383(4), 401(3), 411(7), 519(9)	268, 358									
20.22	556.1479	556.1481	-0.72	C24H29O12NS	516(2), 533(17), 534(4)										
20.67	556.1481	556.1483	-0.43	C24H29O12NS	273(100), 274(15), 315(2), 394(2), 435(55), 436(10)	286									
20.96	556.1481	556.1483	-0.35	C24H29O12NS	273(100), 274(17), 284(9), 297(1), 394(49), 399(11), 417(2), 435(14)	286									
21.26	556.1476	556.1483	-0.06	C24H29O12NS	273(100), 274(11), 315(2), 394(3), 435(65), 436(10)	3287									
33.12	567.1711	567.1708	0.54	C26H30O14	273(100), 315(4), 399(3), 417(11), 435(19)	366									
19.97	572.1430	572.1432	-0.48	C24H29O13NS	289(100), 290(14), 451(12)	292									
33	581.1864	581.1865	-0.14	C27H32O14	273(100), 314(2), 315(4), 383(4), 401(5), 417(7), 419(13), 435(24), 527(4), 561(5)	370									
31.28	595.1660	595.1657	0.34	C27H30O15	287(100), 433(2), 449(25)	266, 348									
28.53	611.1608	611.1607	0.29	C27H30O16	303(100), 397(11), 448(3), 449(1), 465(27)	256, 354									
20.93	611.1605	611.1607	-0.32	C27H30O16	287(29), 365(5), 368(4), 397(35), 436(4), 448(56), 449(100), 450(10), 577(8)	268, 330									
23.3	614.2082	614.2080	0.40	C27H30O15N	273(100), 435(66), 436(4), 597(7)	286									
18.8	627.1555	627.1556	-0.12	C27H30O17	303(24), 465(100), 466(9)	258, 356									
24.56	630.2032	630.2032	0.52	C27H36O16N	289(49), 451(100), 452(3), 586(5), 612(8), 613(29)	284									
24.61	642.1517	642.1512	0.70	C23H31O20N	273(57), 274(8), 315(6), 339(2), 381(3), 485(4), 503(8), 504(3), 521(100), 522(21)	284									
35.34	683.1816	683.1818	-0.28	C30H34O18	273(12), 297(3), 315(7), 381(4), 485(6), 503(28), 521(100), 665(4)	280									
25.92	743.2026	743.2029	-0.44	C32H38O20	303(100), 435(2), 449(3), 465(13), 597(6), 611(38)	256, 354									
28.03	727.2074	727.2080	-0.83	C32H38O19	287(100), 419(3), 433(4), 449(23), 581(6), 595(57)	266, 352									
20.65	757.2185	757.2186	-0.09	C33H40O20	287(48), 433(4), 449(100), 595(36), 611(66), 612(2)	226, 354									
18.64	773.2133	773.2135	-0.24	C33H40O21	303(58), 449(7), 465(100), 611(36), 627(59)	258, 356									
23.9	773.2135	773.2135	0.02	C33H40O21	303(66), 431(4), 449(21), 465(100), 610(55), 611(48), 754(3)	284									
40.47	811.2659	811.2655	0.50	C37H46O20	287(100), 345(5), 433(4), 449(11), 503(2), 595(5), 665(6), 666(2), 680(5), 737(11)	weak									
35.83	813.2444	813.2448	-0.53	C36H44O21	303(100), 331(4), 449(4), 465(9), 505(3), 611(5), 667(13), 680(15), 681(5), 723(4), 737(32), 752(5), 789(7), 794(3), 802(4), 803(26)	weak									
38.33	827.2604	827.2604	-0.04	C37H46O21	303(100), 304(11), 327(5), 345(10), 369(5), 383(4), 401(3), 411(7), 515(9), 615(2), 635(9), 634(4)	256, 356									
38.78	831.2341	831.2342	-0.15	C39H42O20	287(100), 288(3), 365(4), 383(4), 433(3), 449(12), 523(3), 595(6), 669(3), 685(5)	290, 336									
24.39	843.2186	843.2190	-0.43	C36H42O23	287(12), 535(100), 681(5), 697(10)	284									
36.66	847.2289	847.2291	-0.43	C39H44O21	303(100), 30492, 365(3), 383(3), 449(4), 465(11), 539(4), 563(2), 581(3), 611(7), 683(2), 685(6), 701(11), 801(2), 811(2), 822(2), 828(2), 829(2), 830(2)	268, 352									
35.55	873.2443	873.2448	-0.56	C41H44O21	287(100), 389(5), 407(4), 425(16), 433(4), 441(2), 449(21), 565(4), 579(3), 595(7), 711(4), 726(3), 727(7)	284, 370									
33.59	889.2389	889.2397	-1.01	C41H44O22	303(100), 345(4), 389(3), 407(4), 425(13), 449(7), 465(25), 581(7), 611(8), 593(4), 727(7), 743(13)	288, 394									
35.24	903.2552	903.2553	-0.17	C42H46O22	287(100), 291(1), 309(4), 329(4), 419(5), 433(3), 437(2), 449(31), 454(6), 455(14), 471(4), 595(12), 705(2), 721(4), 741(4), 757(9), 883(5), 884(7), 885(3)	328									
17.1	905.2551	905.2557	-0.71	C38H48O25	303(52), 308(20), 309(3), 345(3), 435(10), 449(8), 465(100), 470(3), 580(5), 597(14), 611(37), 613(4), 627(39), 742(35), 743(16), 759(6), 773(87), 775(4)	weak									
33.37	919.2495	919.2503	-0.83	C42H46O23	303(100), 309(4), 345(3), 419(4), 449(5), 455(12), 465(30), 471(4), 611(14), 755(4), 756(4), 757(6), 772(2), 773(9), 901(3)	290, 348									
35.71	923.2822	923.2816	0.66	C42H51O23	303(100), 345(3), 423(3), 441(3), 449(5), 456(9), 465(24), 611(4), 615(6), 761(4), 777(8)	282, 384									
33.98	933.2651	933.2659	-0.89	C43H48O23	287(100), 303(4), 321(16), 339(7), 353(3), 433(3), 449(40), 485(21), 501(3), 595(6), 625(5), 774(7), 787(14)	284, 336									
32	949.2604	949.2608	-0.45	C43H48O24	75(3), 303(100), 321(13), 339(5), 345(3), 449(11), 465(28), 485(16), 501(3), 611(8), 641(4), 787(8), 803(14), 870(2), 887(27), 888(8), 929(3)	332									
34.72	951.3120	951.3129	-0.91	C44H54O23	287(100), 303(3), 321(5), 339(4), 357(5), 433(5), 449(37), 467(6), 485(13), 499(6), 503(18), 517(5), 595(8), 643(7), 647(3), 789(3), 805(9), 932(3)	282, 378									
37.78	955.2550	955.2503	0.85	C45H46O23	273(24), 435(100), 521(29), 907(6), 908(16), 909(4)	366									
32.95	967.3139	967.3137	0.26	C37H58O29	303(100), 321(2), 339(3), 357(4), 449(7), 465(25), 467(4), 485(10), 502(3), 503(12), 611(7), 659(5), 805(4), 821(9)	weak									
17.4	977.3071	977.3074	-0.26	C49H52O21	303(5), 465(18), 611(12), 627(11), 773(100)	270, 354									

58 flavonoids

**More
abundant in
the peel than
in the flesh.**

Iijima et al. (2008)
Plant J.

Annotation for major secondary metabolite Glycoalkaloids

90 glycoalkaloids

Mostly unknown.

**During ripening,
larger compounds
increased.**

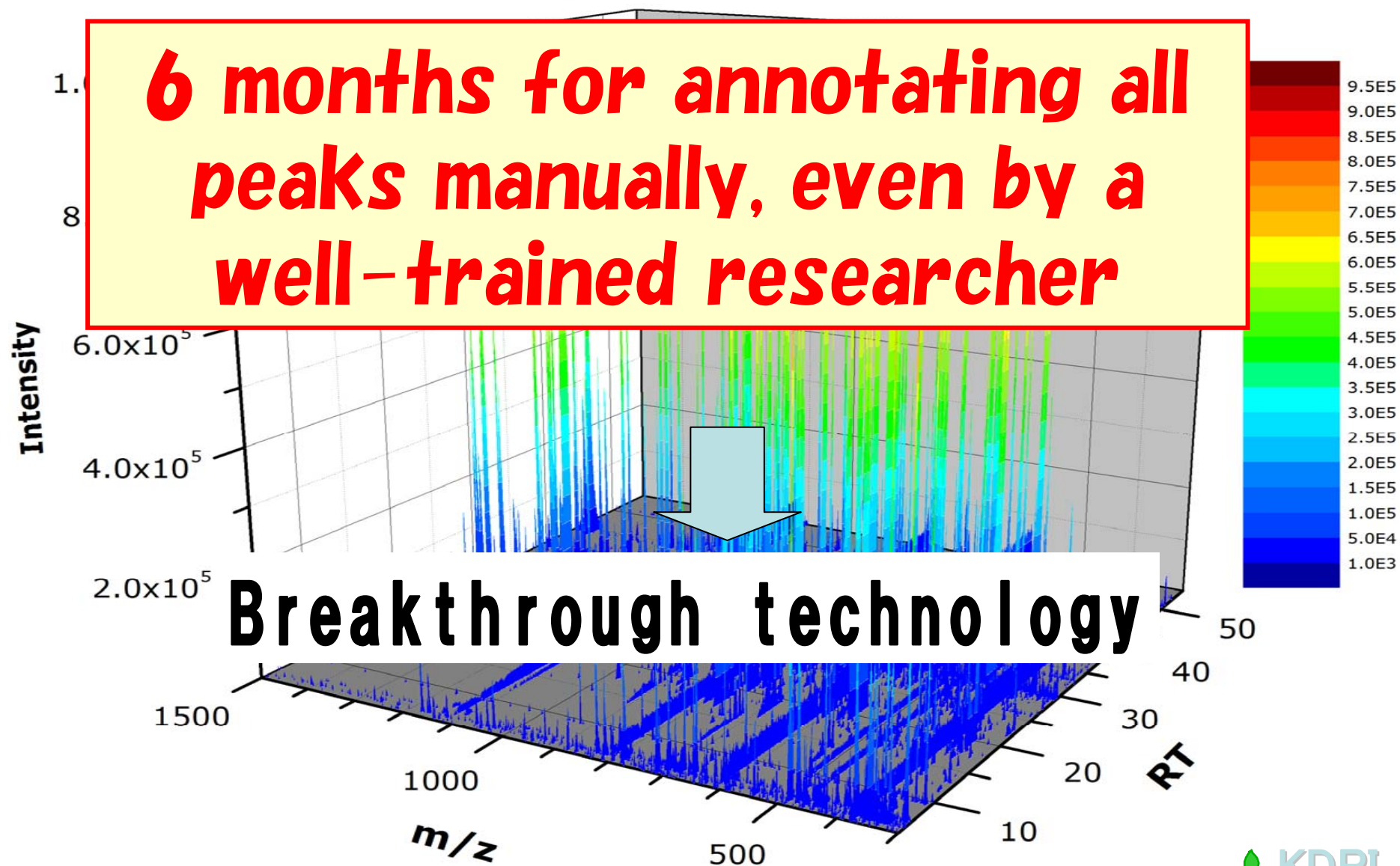
**Further
glycosylation
hydroxylation
acetylation of pre-
existing alkaloids?**

Iijima et al. (2008) Plant J.

	R1	Detected	DEFECT	Formula	G	Y	O	R	L	S	F	MS/MS
1	32.81	738.44257	738.442	C39H63O12N								414(1), 558(4), 576(60), 577(21), 600(4), 704(2), 720(100), 721(42)
2	35.03	738.44275		C39H64O12N								255(49), 273(9), 386(6), 396(40), 414(100), 576(22)
3	31.64	756.45313	756.453	C39H65O13N								738(100)
4	33.4	756.4539		C39H65O13N								722(59), 720(100), 576(66), 558(5), 414(10), 255(6)
5	43.17	842.49579	842.496	C41H69O13N								386(3), 414(8), 456(8), 542(17), 580(70), 720(43), 824(100), 825(54)
6	38.78	872.50322	872.503	C41H71O13N								414(1), 532(16), 576(2), 101(2), 854(100)
7	33.77	906.51062	-0.55	C44H75O18N								415(2), 577(100), 709(2), 739(8), 871(5)
8	42.51	912.47495	912.474	C49H90O18N								394(9), 412(18), 454(40), 540(11), 738(27), 750(69), 890(18), 892(100)
9	38.09	914.39823		C49H90O18N								386(3), 414(2), 456(10), 542(18), 580(40), 720(54), 824(100), 825(66)
10	38.95	914.49188	914.496	C49H90O18N								396(32), 412(12), 414(13), 542(16), 720(49), 734(6), 750(6), 892(6), 894(72), 896(100)
11	43.7	914.49910	914.498	C49H90O18N								414(1), 432(16), 456(16), 542(14), 720(33), 732(7), 894(6), 896(100)
12	29.08	916.49304	916.49	C45H73O18N								417(1), 431(1), 898(100)
13	28.23	918.50993		C45H73O18N								414(4), 432(2), 576(1), 593(1), 898(2), 900(100)
14	29.76	918.50995	918.509	C45H73O18N								410(2), 412(3), 432(3), 576(1), 593(1), 899(1), 900(100)
15	30.13	918.50997	918.506	C45H73O18N								414(2), 432(4), 576(2), 900(100)
16	31.65	918.50999		C45H73O18N								414(1), 432(5), 576(2), 594(2), 898(1), 900(100)
17	32.1	918.50635	918.506	C45H75O18N								396(2), 414(5), 432(3), 576(2), 594(1), 738(2), 756(2), 882(12), 900(100), 901(53)
18	41.03	918.50989		C45H75O18N								414(2), 432(7), 576(1), 594(1), 898(9), 900(100)
19	32.81	920.52319	920.523	C45H75O18N								398(3), 416(6), 578(2), 740(1), 884(100)
20	42.89	940.47127	940.469	C50H87O19N								392(100), 410(4), 578(1), 740(1), 761(3), 899(100), 908(16), 910(42), 925(40)
21	39.15	940.54965	-0.64	C54H81O19N								289(13), 325(7), 365(3), 383(4), 401(14), 419(23), 487(4), 596(36), 599(100), 760(17), 761(32)
22	42.23	942.48971	942.489	C54H81O19N								398(100), 414(100), 531(12), 543(1), 788(12), 812(11), 921(10)
23	34.76	944.51595	944.5	C50H74O19N								396(14), 414(4), 542(20), 720(2), 782(6), 896(18), 914(80), 926(100)
24	40.72	944.80208	944.8	C50H74O19N								396(17), 414(43), 542(8), 718(18), 720(36), 721(17), 780(8), 782(3), 896(16), 912(19), 914(71), 926(100), 927(55)
25	39.34	944.80939		C50H74O19N								890(20), 927(20), 945(73), 946(100), 950(100), 958(2), 957(65), 956(6)
26	31.5	974.49987	974.496	C47H75O20N								410(2), 470(1), 867(3), 896(4), 914(100), 956(6)
27	32.23	976.51411	976.513	C47H75O20N								396(100), 414(100), 516(100), 916(100)
28	37.97	1004.54169	-0.13	C49H81O20N								396(6), 416(11), 578(5), 842(4), 842(7), 986(100)

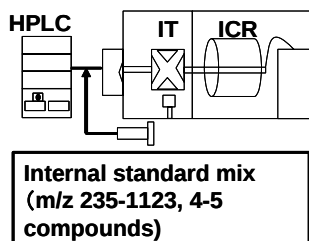
The major problem of this approach

6 months for annotating all peaks manually, even by a well-trained researcher

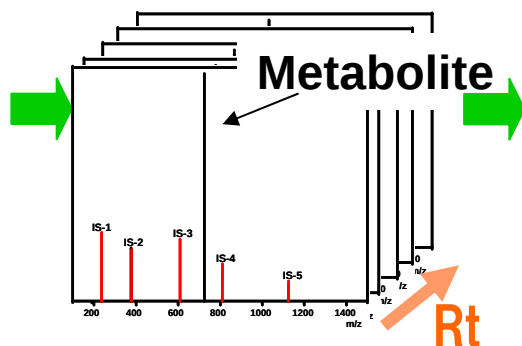


PowerFT: high-throughput pipeline

LC-FTICR/MS analysis



Raw data acquisition



m/z calibration by I.S.

44	485.1030920194518	15618.5859375
45	505.3808537090883	9080.4539609375
46	563.7856319206811	11845.8623046875
47	589.7864132550492	11762.7734375
48	580.1482556075436	8638.587890625
49	580.1803547001539	156112.015825
50	581.183587538724	31624.013671875
51	599.1977307919923	50878.32421875
52	600.1994377527379	12416.8328125
53	609.2806572 23903	113140625
54	610.2849098306476	13073.103515625
55	611.1338920898123	18487.01171875
56	611.1387137719458	23682.087890625
57	611.1460988800231	82982.609375
58	611.1607470282763	1279004.875
59	612.1497411711599	25184.785625
60	612.1846334552172	375310.4375
61	613.186994033382	55208.76953125
62	614.189588747351	9905.435546875

Extract m/z peak groups

Detection of isotopic peak pairs

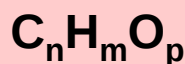
MF calculation

Manual assignment

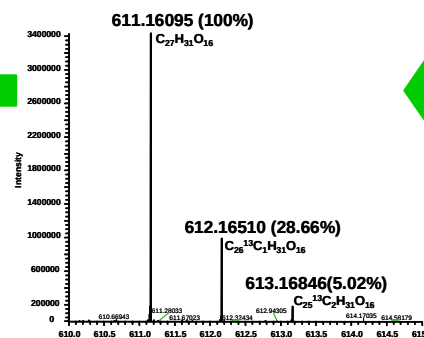
Metabolite annotations

- MSⁿ
- PDA
- Database hit
- Literature

MF confirmation



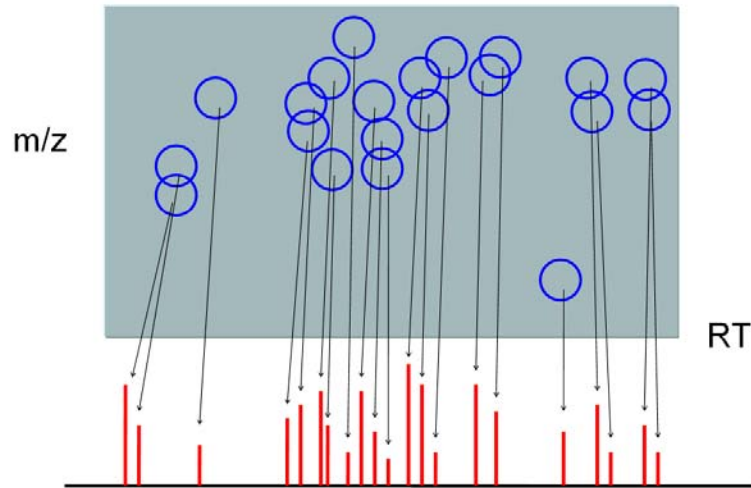
Narrow down candidate formulas by isotopic pattern



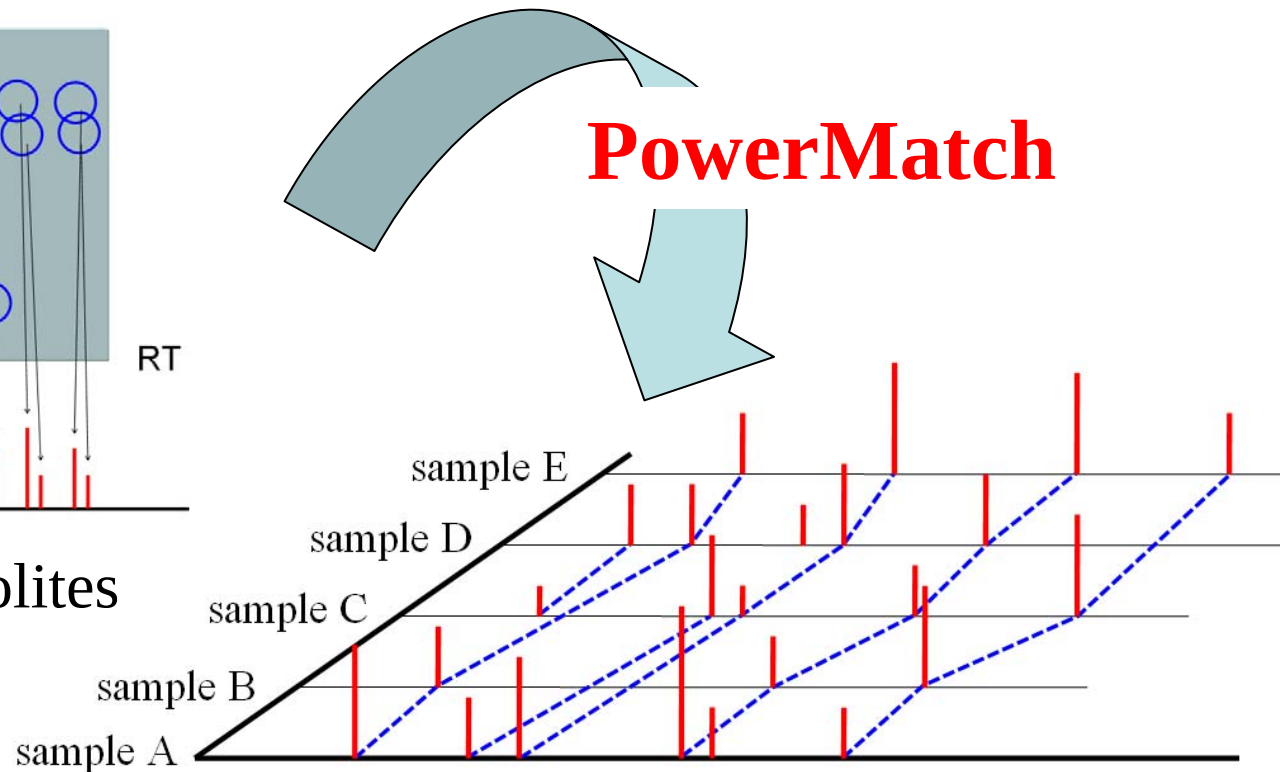
303	611.1607470282763	1279004.875
304	611.1607470282763	1279004.875
305	Total=76	Actual=610.1534718668763
306	C27H34O16	0 0 0 0 0 0 0 610.153384 8.708257632861205E-5
307	C25H18N14O8	0 0 0 0 0 0 0 610.153374 9.758477631294227E-5
308	C10H14N2O7	0 0 0 0 0 0 0 610.153376 4.049987237522148E-4
309	C12H26N12O17	0 0 0 0 0 0 0 610.153887 4.155008297385448E-4
310	C40H22N2O5	0 0 0 0 0 0 0 610.152871 6.001882762644123E-4
311	C41H18N6O1	0 0 0 0 0 0 0 610.154209 7.372445297478042E-4
312	C11H10N8O21	0 0 0 0 0 0 0 610.152550 9.219118762757718E-4
313	C9H18N2O11	0 0 0 0 0 0 0 610.152539 9.324140762601019E-4
314	C7H6N3O1	0 0 0 0 0 0 0 610.152529 9.429182762444921E-4
315	C26H14N18O2	0 0 0 0 0 0 0 610.154711 0.001238828023585879
316	C28H22N14O12	0 0 0 0 0 0 0 610.154722 0.00125030302236897044
317	C24H22N10O10	0 0 0 0 0 0 0 610.152036 0.0014349875763252587
318	C22H18N2O4	0 0 0 0 0 0 0 610.152026 0.001445498776308583
319	C11H10N8O3	0 0 0 0 0 0 0 610.156214 0.0017424115236508442
320	C13H22N16O13	0 0 0 0 0 0 0 610.156224 0.001752913727488812
321	C15H6N4O23	0 0 0 0 0 0 0 610.156235 0.001763415927931914
322	C8H10N3O5	0 0 0 0 0 0 0 610.151202 0.002258268267274183
323	C6H10N3O5	0 0 0 0 0 0 0 610.151191 0.0022803290762567495
324	C23H22N8O8	0 0 0 0 0 0 0 610.156059 0.0025877430236968021
325	C23H22N8O14	0 0 0 0 0 0 0 610.156069 0.002772410376337575
326	C21H14N2O4	0 0 0 0 0 0 0 610.156889 0.0027829125763219054

How to compare metabolomes

2-D image of LC-FT/ICR-MS



1-D image of metabolites



Alignment of metabolites with information of Retention time, mass values, MS/MS patterns on 1-D

PowerMatch: alignment software

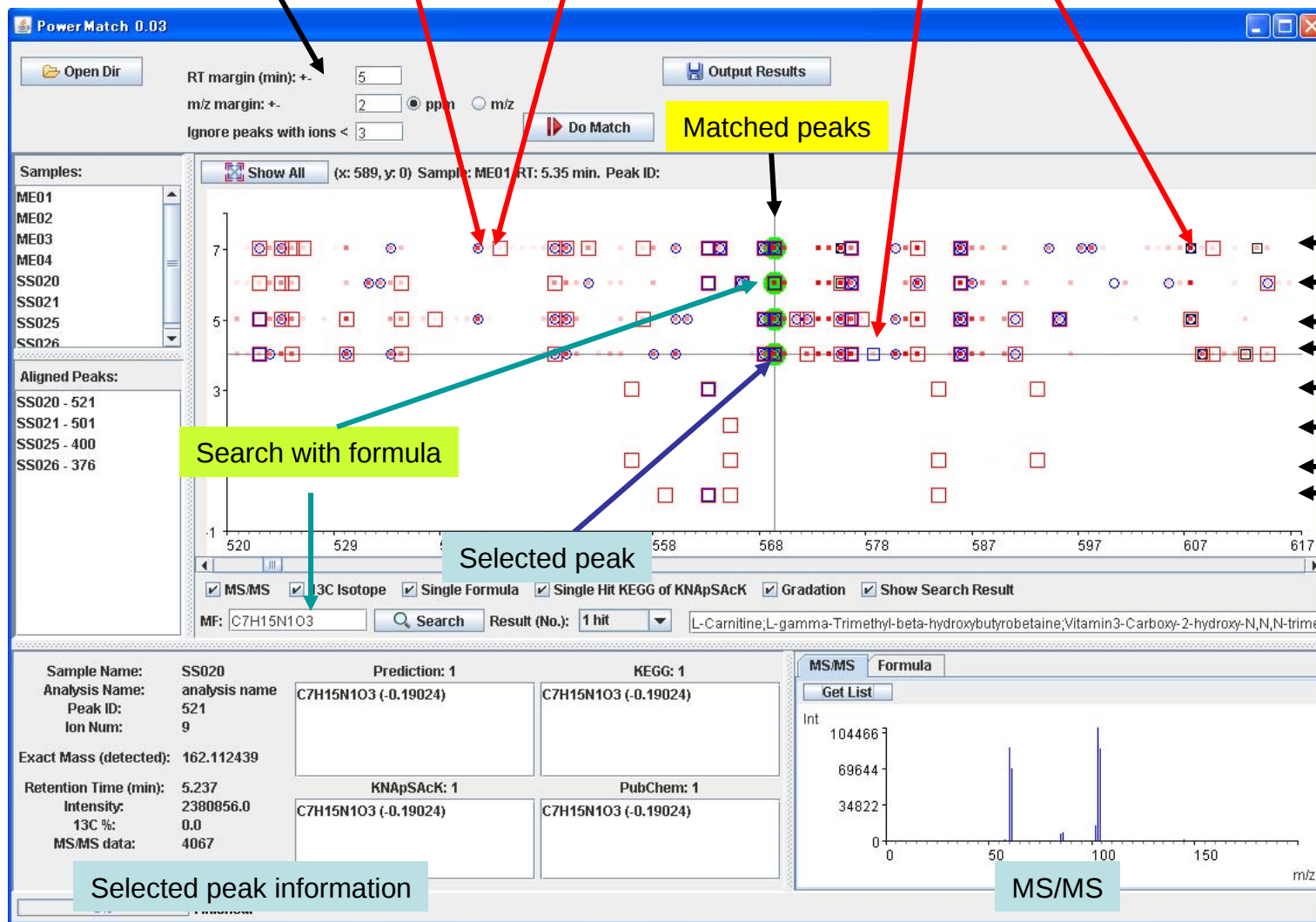
○: peak accompanied by MS/MS data

□: peak showing single formula in KNApSack/KEGG

parameters

□: peak showing single formula

□: peak accompanied with ^{13}C isotope peak



samples

Links to Metabolite Databases

The image displays four windows of metabolite databases, illustrating their interlinking capabilities:

- JustMatch:** A window showing a list of samples on the left and a detailed view of compound C10051 on the right. The compound details include its name (Gossypin; Gossypetin 8-O-glucoside), formula (C₂₁H₂₀O₁₃), mass (480.0904), and chemical structure.
- KEGG:** A window showing the KEGG Compound page for C10051, providing similar information to JustMatch, including the chemical structure and a comment that it is a flavonol 8-glycoside.
- PubChem:** A window showing the PubChem Compound Summary for CID 5281621, which is Gossypin. It includes the chemical structure and links to related compounds and substances.
- KnapSack:** A window showing the KnapSack Metabolite Information page for Gossypin. It includes a table of metabolite information and a table of MS/MS data.

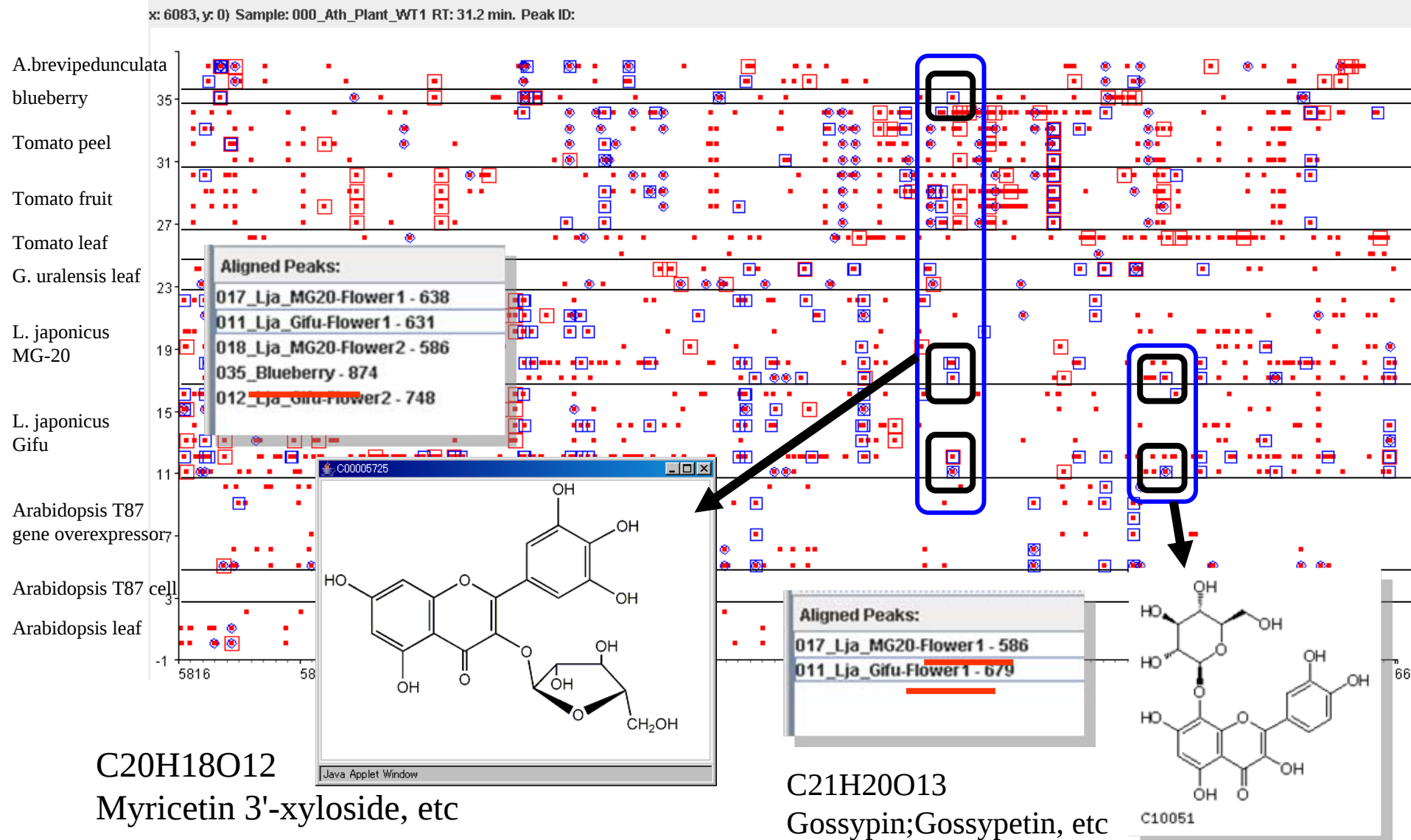
Arrows indicate cross-linking between the databases:

- An arrow points from the KnapSack window to the KEGG window.
- An arrow points from the PubChem window to the KnapSack window.

A red box highlights a table of MS/MS data in the KnapSack window, showing the following information:

DB	M.F.	Mass	dPPM	ID	Name
KEGG	C21H20O13	480.090390...	-0.53509	C10051	Gossypin;Go...
KNAPSAcK	C21H20O13	480.090390...	-0.53509	C21H20O13...	Myricetin 3'-g...
KNAPSAcK	C21H20O13	480.090390...	-0.53509	C21H20O13...	Zeravschan...
KNAPSAcK	C21H20O13	480.090390...	-0.53509	C21H20O13...	Gossypetin...
KNAPSAcK	C21H20O13	480.090390...	-0.53509	C21H20O13...	6-Hydroxytri...
KNAPSAcK	C21H20O13	480.090390...	-0.53509	C21H20O13...	Myricetin 3-gl...
KNAPSAcK	C21H20O13	480.090390...	-0.53509	C21H20O13...	Gossypetin...
KNAPSAcK	C21H20O13	480.090390...	-0.53509	C21H20O13...	Tagetelin
KNAPSAcK	C21H20O13	480.090390...	-0.53509	C21H20O13...	Gossypetin...

Comparison of 38 plant metabolomes



Metabolite annotation database KOMICS

Kazusa Omics Database

KOMICS

Login

HOME Browse Search Download Help

What's New
April 2, 2008
Two samples of *C. elegans* were added

Statistics
Species: 8
Samples: 562
Identified Metabolites: 271
Total Metabolite Peaks: 13962

Links
Kazusa Metabolome Home
KaPPA-View
MassBase
KNAPSAcK
KEGG
PRIME
Metabolome.jp
MassBank

Contact us

Overview
Comprehensive analysis of metabolites in organisms are compiled in this database. Metabolite peaks detected by various analytical instruments are deposited with their annotation. KOMICS provides browsing, searching and comparing the metabolites across species and experimental conditions, and that facilitates to understand and utilize the chemical diversity of the living organisms for biomarker discoveries, gene resource discoveries and material productions.

Plants	-		<i>Arabidopsis thaliana</i>																			
	-		Tomato																			
	-		Rice																			
Microbes	-		<i>E. coli</i>																			
	-		Yeast																			
Animals	-		<i>C. elegans</i>																			
	-		Zebra fish																			
	-		Human																			

Species View (Sample Image)

<http://webs2.kazusa.or.jp/komics/>

