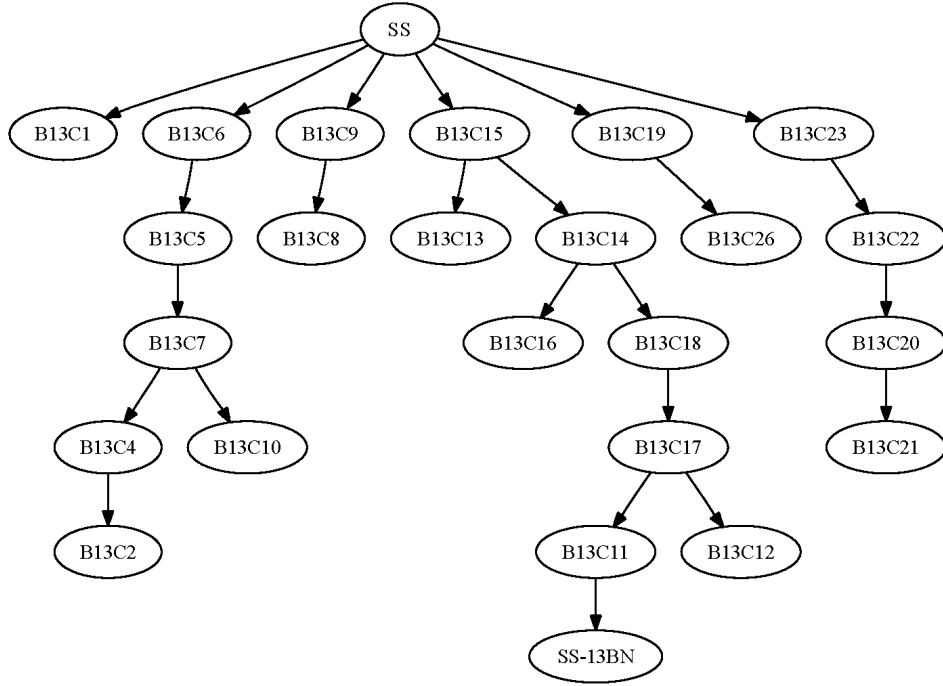


Suppl. figure 1. MST tree for the MAP study in the rat.



Suppl. table 1. Pseudo-code for a Minimum Spanning Tree based on Kruskal's algorithm.

Because each vertex is in exactly one tree, we use a collection of disjoint sets as data structure with the following functions:

make\_set(x): create a disjoint set with element x.

find(x): find which set is element x in

union(x,y): replace set containing x and y with their union

Given graph  $G=(V,E)$ , let T be the MST under construction

1. Sort edges  $E: (u,v)$  by cost/weight
2.  $T = \{\}$
3. For x in V  
    make\_set(x)
4. For  $(u,v)$  in E (increasing order):  
    If  $\text{find}(u) \neq \text{find}(v)$   
         $T \leftarrow T \cup \{(u,v)\}$   
        union(u,v)  
    end if  
end For

Suppl. table 2A. Mean arterial pressure (MAP) in females. QTL locations highlighted with a black border were identified with the common segment method (Moreno et al. 2007), whereas QTLs highlighted in pink were identified with the sequential method.

marker	D13Rat7	D13Rat60	D13Rat11	D13Rat15	D13Rat20	D13Rat88	D13Rat12	D13Got22	D13Rat91	D13Rat101	D13Rat127	D13Rat176	D13Rat178	D13Rat177	D13Rat151	D13Rat61	D13Got45	D13Got51	D13Rat46	D13Rat191	D13Rat18	D13Rat57	D13Rat82
SS	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C1	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B
B13C2	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B
B13C4	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B
B13C5	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C6	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C7	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C8	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C9	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C10	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C11	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C12	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C13	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C14	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C15	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C16	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C17	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C18	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C19	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C20	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C21	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C22	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
B13C23	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
SS-13BN	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B	B

vs	otfl	sig	pval	pval/nTest	str1	str2	sig	pval	pval/nTest
B13C1		1	0.00062	0.000396	SS	B13C1	1	0.00062	0.000396
B13C2	0	0.032583	2.246146		B13C4	B13C2	0	0.005429	0.100269
B13C4	1	3.81E-05	0.000915		B13C7	B13C4	0	0.736689	17.88005
B13C5	1	0.00128	0.03072		B13C6	B13C5	1	0.000291	0.00638
B13C6	0	0.38541	9.249848		SS	B13C6	0	0.38541	9.249848
B13C7	1	0.000495	0.01872		B13C5	B13C7	0	0.931873	22.36496
B13C8	0	0.003082	0.073966		B13C9	B13C8	0	0.554823	13.31814
B13C9	0	0.018623	0.446362		SS	B13C9	0	0.018623	0.446362
B13C10	1	0.001819	0.043664		B13C7	B13C10	0	0.568458	13.64298
B13C11	1	7.5E-05	0.001801		B13C17	B13C11	0	0.246361	5.912655
B13C12	0	0.284472	6.82734		B13C17	B13C12	0	0.007208	0.172992
B13C13	0	0.531298	12.75116		B13C15	B13C13	0	0.470223	11.28536
B13C14	0	0.607879	14.58909		B13C15	B13C14	0	0.07413	1.779118
B13C15	0	0.224085	5.378035		SS	B13C15	0	0.224085	5.378035
B13C16	0	0.27833	6.679329		B13C14	B13C16	0	0.091775	2.202612
B13C17	1	0.00046	0.01041		B13C18	B13C17	0	0.302368	7.258821
B13C18	1	0.000606	0.014534		B13C14	B13C18	1	4.41E-05	0.001053
B13C26	1	0.000114	0.002725		B13C19	B13C26	0	0.005454	0.130889
B13C19	0	0.205545	4.933086		SS	B13C19	0	0.205545	4.933086
B13C20	0	0.197381	4.737153		B13C22	B13C20	0	0.809791	19.43499
B13C21	0	0.196364	4.712735		B13C20	B13C21	0	0.784141	18.81939
B13C22	0	0.368072	8.833728		B13C23	B13C22	0	0.653561	15.68546
B13C23	0	0.518482	12.44356		SS	B13C23	0	0.518482	12.44356
SS-13BN	1	5.93E-10	1.42E-08		B13C11	SS-13BN	0	0.343175	8.236204

Black Bordered boxes were identified by the paper  
 Pink boxes were identified by the sequential method



Suppl. table 2C. Genetic markers for studying MAP (see Moreno et al. 2007 for details).

<b>Marker</b>	<b>Left</b>	<b>Right</b>
D13Rat7	1084269	1084470
D13Rat60	13053809	13053933
D13Rat111	32030140	32.30259
D13Rat115	35719011	35719148
D13Rat20	38329090	38329229
D13Rat88	42627611	42627837
D13Rat123	44872182	44872315
D13Got22	45828609	45828716
D13Rat91	46911976	46912163
D13Rat101	49428398	49428577
D13Rat127	55373162	55373537
D13Rat178	59830974	59831216
D13Rat179	60669452	60669696
D13Rat77	63175275	63175600
D13Rat151	64083141	64083335
D13Rat61	68034721	68034927
D13Got45	69494048	69494326
D13Got51	72638817	72638941
D13Rat46	77336099	77336322
D13Rat197	80109433	80109632
D13Rat183	80966179	80966380
D13Rat57	84280243	84280390
D13Rat192	100133218	100133413



Suppl. table 3. SNP markers for Chromosome 13.

<b>Marker</b>	<b>Left</b>	<b>Right</b>
D13M236	4363272	4363368
D13M218	19636983	19637157
D13M88	38770728	38770896
D13M99	77262943	77263147
D13M126	85806862	85807000
D13M106	93838592	93838725
D13M77	118025687	118025961