Supplementary data for selected publications

Primdahl H., Wikman FP., von der Maase H., Zhou X, Wolf H, Orntoft TF.: Allelic Imbalances in Human Bladder Cancer: Genome-wide Detection With High-Density Single-Nucleotide Polymorphism Arrays.

Journal of the National Cancer Institute 2002 Feb 94(3) 216-223.

PubMed

Loci with allelic imbalance in >=3 tumors within less than 16MB nucleotide distance

Chromosome	Position (MB)[a]	T1[b]	T2-4[c]	Number of tumors	patients
1	133.2	2	2	4	2
2	212.7-220.9	0	4	4	4
6	11.1-22.3	3	4	7	5
6	40.0-40.8	1	2	3	2
6	114.0-118.4	1	2	3	2
6	158.4-165.2	2	4	6	4
7	31.4-40.0	1	3	4	3
8	20.2-28.2	3	2	5	3
8	80.3-92.8	2	2	4	3
8	118.6-127.5	1	2	3	2
9	3.1-7.5	1	3	4	3
9	10.6-22.5	1	2	3	3
9	40.0-48.0	1	6	7	6
9	70.5-82.1	2	4	6	4
9	89.3-93.6	0	3	3	3
9	102.1-106.6	3	4	7	4
11	11.1	1	4	5	4
11	19.6	2	1	3	2
11	67.2-67.8	1	2	3	2
11	101.3-103.8	1	2	3	2
12	77.2-88.6	2	2	4	3
13	37.0-44.6	1	2	3	3
13	73.7-76.7	1	2	3	2
13	85.8-95.2	2	2	4	3
14	35.8-36.5	1	2	3	2
14	67.9-76.3	1	2	3	2
14	92	1	2	3	2
16	8.0-15.2	1	2	3	2
17	0.9-3.8	3	3	6	4
17	15.0-20.9	2	2	4	2
17	30.9-33.0	1	2	3	2
18	56.2-71.8	2	3	5	4

[a]Distance in megabases from the p-end of the affected chromosome.

[b]Number of T1 tumors with allelic imbalance in this interval, c Number of T2-4 tumors with allelic imbalance in this interval, Total number of tumors with allelic imbalance in this interval.

[c]Number of patients with allelic imbalance in this interval.