

Table 1. Restriction landmark genomic scanning (RLGS) analysis along 6q23-24 in primary HNSCC and NSCLC

NSCLC	n2D61	a5G27	a6E12	n2E20	a2F23	a5E34	n4F51	a5F58	n6E07	n3G47
1			■							
2			■	■	■	■				
3			■	■						
4	■		■				■		■	■
5			■	■		■				
6										
7			■			■				
8			■	■		■				■
9			■							
10			■				N			
11								N		■
12		N	■					N		
13								N	N	
14			■	■		■		N	N	
15			■					N		
16			■	■			■	N		
17			■	■		■	N			■
18			■			■				
19			■			■				
20			■				■			
21			■							
22	■		N	■						
23	N	N	■	■						■
24			N	N		N				

HNSCC	n2D61	a5G27	a6E12	n2E20	a2F23	a5E34	n4F51	a5F58	n6E07	n3G47
1										■
2			■	■		■				
3			■			■				■
4			■							
5			■			■				
6			■							
7			■			■				
8			■							
9										
10										
11			■							
12		N	■	■		■				
13										
14										
15			■							

RLGS analyses in individual samples are listed in rows, and RLGS fragments are listed in columns. Methylated (filled box), unmethylated (open box), and partially methylated

(shaded box) NotI (n) and AscI sites (a) are listed with the RLGS master profile addresses. N refers to a RLGS fragment that cannot be analyzed.