

Errata

Volume 35, No. 3, page 1271: The name of the fifth Author should read:

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Volume 35, No. 5, page 2609: Table V should be replaced by: (*see Table V on the next page*)

Volume 35, No. 11, page 6050: The third line of the right column should read:

lung injury (ALI) mice (38, 39). Radiation-induced muscle

Location	Type	PolyPhen		Gene		MUTYH	MTH1	MHH	MUH1	FAM175A	XRCC2	PM52		NBN	ATM		BRCA2		RAD51D	BRCA1		BRIP1	TP53				
		SNP	SNP	SNP	SNP							SNP	SNP		SNP	SNP	SNP	SNP		SNP	SNP			SNP	SNP	SNP	SNP
Canada	# of patient (n=10)	4	2	4	0	0	7	6	3	7	8	2	10	3	2	2	10	6	10	2	2	8	7	3	7	8	
		total # of alleles (n=20)	4	2	4	0	0	14	12	6	14	16	4	20	6	4	20	12	20	4	4	16	14	7	10	11	
Saudi Arabian	# of patient (n=10)	20	10	25	0	0	45	40	15	40	50	10	90	15	10	10	35	100	10	10	40	35	15	15	50	55	
		total # of alleles (n=20)	4	0	9	2	8	8	6	3	2	5	1	9	1	5	5	10	5	5	3	6	6	6	6	9	5
Saudi Arabian	# of patient (n=20)	6	0	9	2	10	8	8	3	2	6	1	16	1	5	2	20	9	20	5	3	7	7	6	1	11	7
		total # of alleles (n=40)	30	0	45	10	40	40	15	10	30	5	80	5	25	10	40	45	100	25	15	35	35	30	5	55	35
Total	# of patient (n=20)	8	2	13	2	15	12	12	6	9	13	3	19	4	7	7	20	11	20	7	5	14	13	4	16	13	
		total # of alleles (n=40)	10	2	14	2	19	14	16	6	10	16	3	34	4	13	40	16	40	7	5	15	14	4	21	18	
Global	Global MAF	25	5	35	5	47.5	35	40	15	25	40	7.5	85	10	37.5	17.5	40	100	17.5	12.5	37.5	35	32.5	10	52.5	45	
		allele frequency (%)	G=0.3135/1570	A=0.101/21	T=0.3662/1834	NA	G=0.4593/2300	T=0.3339/1672	A=0.193/421	G=0.172/374	T=0.423/921	T=0.0525/263	T=0.1168/585	T=0.1120/561	A=0.3582/1794	T=0.070/153	G=0.3570/1788	T=0.0242/121	C=0.1508/755	T=0.0950/476	C=0.3277/13	C=0.3526/1766	C=0.303/661	G=0.4561/2284	T=0.035/168	A=0.3722/1864	G=0.398/867
Global	allele frequency (%)	31.35	1	36.6	NA	45.9	54.1	65.6	19.3	17.2	42.3	5.25	88.3	11.2	35.8	7	35.7	97.6	15	9.5	32.7	35.3	30.3	54.5	3.35	62.8	60.2
		allele frequency (%)	G=0.3135/1570	A=0.101/21	T=0.3662/1834	NA	G=0.4593/2300	T=0.3339/1672	A=0.193/421	G=0.172/374	T=0.423/921	T=0.0525/263	T=0.1168/585	T=0.1120/561	A=0.3582/1794	T=0.070/153	G=0.3570/1788	T=0.0242/121	C=0.1508/755	T=0.0950/476	C=0.3277/13	C=0.3526/1766	C=0.303/661	G=0.4561/2284	T=0.035/168	A=0.3722/1864	G=0.398/867