Characteristics of induced mutations in offspring derived from irradiated mouse spermatogonia and mature oocytes

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Recap – WGS after exposure to lonizing Radiation

- Adewoje
 - Mice irradiated with 3Gy x-rays
 - 6 mice offspring 8 weeks after exposure
 - Found increase in multisite de novo mutations, indels and cnvs
- Pilot Study
 - Soldiers exposed to unknown quantities of IR
 - Increase in MSDNs

Study Setup

- Exposure: 4Gy gamma-rays
- Twofold Setup:
 - a) Irradiated male mice
 - Mating sixteen weeks after exposure
 - b) Irradiated female mice
 - Mating immediately after exposure
- Non-irradiated parents as control cohort
- 6 offspring and controls per experiment



Whole Genome Sequencing

- All mice were Whole genome sequenced
- Variants detected by HaplotypeCaller
- Called in Region (54% of total mouse autosomal genome):
 - 50% < x < 300% coverage compared to peak mapped to autosomal data
 - 80% < ratio of high quality reads
 - Overlap of these regions in all samples
- Classes of mutation:
 - SNV (total 339)
 - Indel (total 79)
 - Multisite Mutation (Length <= 100bp)
- CNVs not included

Filtering Criteria

• SNV:

- AAF < 0.1 in parents
- AAF > 0.25 in case
- Review by manual inspection
- Indel:
 - Excluded:
 - Mutations in mononucleotide repeat sequences > 7 in length
 - Dinucleotide > 4
 - Trinucleotide (or more) > 2
- Validation by Sanger
 - 53/53 SNV, 40/40 Indels

Age Effects

- Estimate parental age effects for mice
- Base reference: de novo SNV count at 8 weeks parental age

	No. of increase of mutations (/ week)	Percentage of increase in the number of mutations per week to the number of spontaneous mutations in F1 of 8 week-old
		parents
Paternal effect	0.46 (=[8.8-5.6]/7) [-0.063, 0.99] ¹⁾	4.5% (= 0.46 / 10.3)
Maternal effect	0.24 (=[6.9-4.7]/9) [-0.18, 0.70]	2.3% (= 0.24 / 10.3)

1) 95% CI was calculated by 1,000 times of simulations based on Poisson distribution and binominal distribution.

SNV Mutation Rates

- Context:
 - No increase in Pilot Study, Adewoje
- After correcting for age effects:
 - No significant increase

Group	No. of mutations ^a	Mutation rate (95% CI)	P ^b	No. of adjust	mutations ed for parental age ^c	P ^d
SNVs						
Spermatogonia exposure						
Before IR	70	$4.5 imes 10^{-9} (3.5 - 5.7 imes 10^{-9})$	_	70		
After IR	128	8.3×10 ⁻⁹ (6.9–9.8×10 ⁻⁹)	0.0001	65.1		0.91
Mature oocyte exposure						
Before IR	54	$3.5 imes 10^{-9} (2.6 - 4.6 imes 10^{-9})$		54		
After IR	84	5.4×10 ⁻⁹ (4.3–6.7×10 ⁻⁹)	0.012	54.6		0.96

Non-SNV Mutations

• Indels and Multisite Mutations are increased after exposure

Group	No. of mutations ^a	Mutation rate (95% CI)	P ^b	No. of adjust	f mutations ted for parental age ^c	P ^d
Indels			1			1
Spermatogonia exposure						
Before IR	5	$3.2 imes 10^{-10} (1.0 - 7.5 imes 10^{-10})$		5		
After IR	42	$2.7 \times 10^{-9} (2.0 - 3.7 \times 10^{-9})$	<0.00001	35.9		<0.00001
Mature oocyte exposure				·		
Before IR	7	4.5 $ imes$ 10 ⁻¹⁰ (1.8–9.3 $ imes$ 10 ⁻¹⁰)		7		
After IR	25	$1.6 \times 10^{-9} (1.0 - 2.4 \times 10^{-9})$	0.0022	22.1		0.0095
Multisite mutations						
Spermatogonia exposure						
Before IR	2	$1.3 imes 10^{-10} (0.16 - 4.6 imes 10^{-10})$		2		
After IR	10	$6.5 imes 10^{-10} (0.31 - 1.2 imes 10^{-9})$	0.031	10		0.031
Mature oocyte exposure						
Before IR	0	0 (0-2.4 × 10 ⁻¹⁰)		0		
After IR	10	6.5 × 10 ⁻¹⁰ (0.31–1.2 × 10 ⁻⁹)	0.0031	10		0.0031

Characteristics of IR induced mutations

- Significantly increased number of Indels
- Two main types of deletions:
 - Single-nucleotide in mononucleotide repeat sequences
 - Deletions (1-35bp) in non-repeat sequences
- Both types appear in higher frequency in offspring of irradiated father

			Length of	No. of indels ^b	
	Experiment	Timing of mating	mononucleotide repeats ^a	Deletion (-1 base)	Insertion (+1 base)
S	Spermatogonia exposure	Before irradiation	Non-repeat	1	1
			2	0	0
			3	0	0
			4-7	0	0
		After irradiation	Non-repeat	2	2
			2	8	0
			3	3	1
			4-7	3	2
	Mature oocyte exposure	Before irradiation	Non-repeat	0	0
			2	0	1
			3	1	0
			4-7	0	2
		After irradiation	Non-repeat	0	1
			2	4	2
			3	1	0
			4-7	1	0

Characteristics of IR induced mutations

• Both Indel classes significantly increased compared to unexposed



Characteristics of IR induced mutations

- Multisite mutations (<100bp)
- "Most" within 15bp
- Total 20 mutations, 8 tandem
- C>A Substitutions increased
 - No multisites characteristic to errorprone polymerase (GC>AA, GA>TT)
 - No C>T at CpG sites



Substitution

Conclusions

- DNA repair mechanisms has difficulty:
 - (micro-)homologous indels
 - Indels in repeat sequences
- DNA repair of double strand breaks, strand breaks at repeat sites
- Multisite mutations result from clustered lesions
 - Potential signature of IR induced mutations
- No phenotype observed

Thank you for your attention!