

Reprocessing of Swabs • Obtain DNA profiles in case of instrument or cartridge/chip failure • 9/155 (5.8%) RapidINTEL™ sample cartridges with no possible sizing; all from the same lot number • Enhance quality of initial DNA profile

Reprocessing of Swabs on RapidHIT ID

- 73.7% (14/19) had higher fluorescence signals compared to their initial profiles
- 63.2% (12/19) had 1 to 20 additional alleles compared to their first pass profiles
- 36.8% (7/19) had > 80% alleles compared to 26.3% (5/19) for the first pass profiles and 57.9% (11/19) ≥ 70% alleles compared to 36.8% (7/19)
- Reprocessing increased the number of DNA profiles eligible to search a national DNA database (+3 profiles for Canada and +4 for the US)

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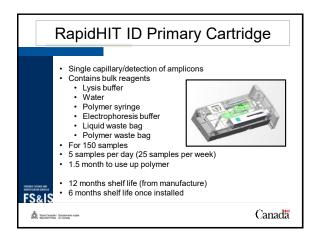
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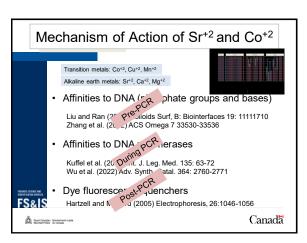
Canada

Canada

Considerations Before Adoption ANDE 6C (ANDE Corporation) I-chip (all steps included) Waste Waste Primary cartridge DNA fragment separation and detection NTEL Sample cartridge Sample lysis, DNA capture, DNA amplification



Gamma Radiation Dose Rates						
http://www.radprocalculator.com/		Distance from	m source (Prima	ary Cartridge): 3	0.5 cm (1 foot)	
		mSv/hr	mSv/hr	mSv/5 min	mSv/5 min	
		Cs-137	Co-60	Cs-137	Co-60	
	1.5 month	→ After 15	Samples	After 25	samples +	1 week
		0.01 M - Level A activity				
Without shielding		18.5	416	0.26	5.8	
Lead blanket (0.635 cm solid lead equivalence)		9.9	272.7	0.13	3.8	
3 cm thick lead		0.4	94.5	0.01	1.3	0.3 μSv/5min
6 cm thick lead	10 μSv/hr	0.01	17.3	0.0003	0.24	
		0.05 M - Level B activity				
Without shielding		92.7	2079.8	1.3	28.9	
Lead blanket (0.635 cm solid lead equivalence)		40.9	1363.3	0.69	18.9	
3 cm thick lead		2	472.6	0.05	6.6	2 μSv/5min
6 cm thick lead	40 μSv/hr	→ 0.04	86.3	0.002	1.2	
		0.10 M - Level C activity				
Without shielding		184.6	4159.6	2.6	57.8	
Lead blanket (0.635 cm solid lead equivalence)		81.4	2726.7	1.4	37.9	
3 cm thick lead		4	945.2	0.1	13.1	3 μSv/5min
6 cm thick lead	90 μSv/hr	→ 0.09	172.5	0.003	2.4	
Gamma radiation dose limit for the general public: 1 mSv/yr (20 μSv/week, 0.5 μSv/hr, 0.042 μSv/5 min)						



Mechanism of Action of Sr ⁺² and Co ⁺²						
PCR reaction mix Primer mix Size Standard L5	Lysis Chamber 300 µL Prep-N-Go Buffer Bubbling agitation, Heat: 10 min/85°C PCR Zone 1.2 mm FTA paper disc < 12 µL/32 cycles, Platinum™ Taq polymerase	ANDE 6C I-chip Swab Chamber 300+ µL Guanidium-based Buffer Chaotic bubbling, no heat Silica membrane/DNA purification Conc. Module 100% lysate				
	oCl ₂ impacts DNA polymera in loss STR allele detection	PCR Zone				

Conclusions for Mock "R" Event

Both Rapid DNA instruments:

- Showed similar first pass success rates (profiles with a minimum of 10 autosomal STR loci)
- Produced complete STR profiles from 1 µL and 10 µL saliva on metal plates or PVC caps in the presence of non-radioactive Cs, Sr and Co salts (Level A/10 mM for CoCl₂ and SrCl₂; Levels A-C 10-50-100 mM for CsCl₂)
- Had profiles adversely impacted using CoCl2 and SrCl2 at Levels B and C; interference at pre-PCR/during PCR
- Showed a similar success in generating DNA profiles eligible to be searched against a national DNA database

Conclusions for Mock "R" Event

- Reprocessing of contaminated samples may increase allele calls and number profiles eligible to be searched against national DNA databases
- The ANDE™ 6C offers quick disposal of all radioactive materials associated with contaminated biological samples into appropriate shielded containers
- The ANDE $^{\mbox{\tiny TM}}$ 6C does not require recalibrations upon deployment and reagent refrigeration
- In a true "R" event, the production of complete Rapid DNA profiles will depend on the amount and type of radiation received by the biological samples. (Level A)

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	Processing biological samples from simulated radiological terrorist events using Rapid DNA instruments: Caustal J. Friegans **, Nancy Lustin* *** *** *** *** *** *** ***			
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	the increase rate Canada			