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Towards an HIV Cure **SIAS** 

Pardons Marion, HIV Cure Research Center (Belgium) Session 3: Cure advances globally

# HIV reactivation from latency using a Tat compound

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## Conflict of interest disclosure

This work was done in collaboration with Janssen and funded by VLAIO



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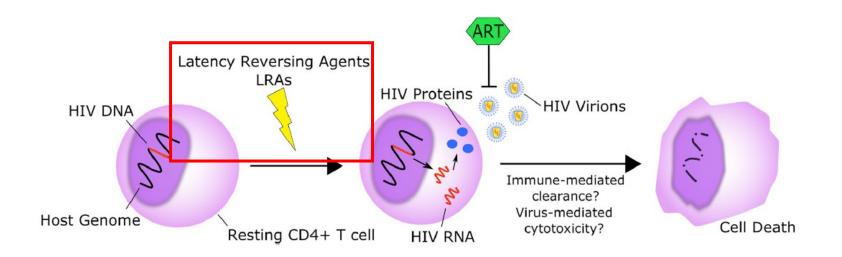
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#### **Shock-and-kill strategy**

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Mitogens	Other classes of LRAs			
PMA, PHA, CD3/CD28	HDACi, PKC agonists, etc			
Gold standard for in vitro assays	Not as potent as mitogens to reactivate HIV			
Highly toxic $ ightarrow$ Not in the clinic	Safe to be used in vivo			
Induces global T cell activation	Some classes do not induce global activation			

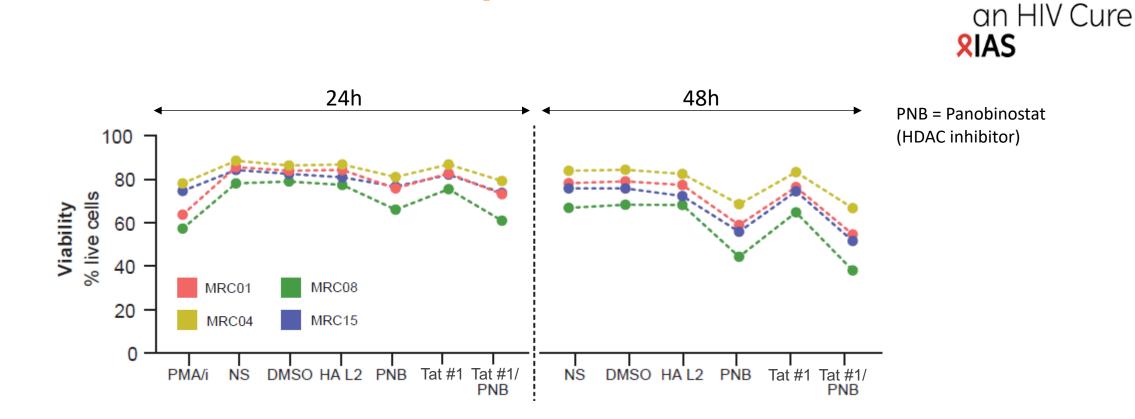
Identifying compounds that reactivate HIV efficiently without modifying the transcriptome/phenotype of the cells is of interest to study the profile of latently infected cells

#### **Clinical characteristics of the participants**

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MRC01         55         M         0.3         44         B         <20		•		004/000						
MRC02         62         M         1.2         484         B         <20	Patient ID	Age	Gender	CD4/CD8 ratio	NADIR	Subtype	VL	Time since infection	Time to ART	ART duration
MRC03         33         M         0.7         114         F1         <20         30.7         16.0         14.6           MRC04         51         M         1.0         171         B         <20	MRC01	55	M	0.3	44	В	<20	NA	NA	18.5
MRC04         51         M         1.0         171         B         <20         NA         NA         18.4           MRC05         42         M         1.3         492         B         <20	MRC02	62	М	1.2	484	В	<20	NA	NA	4.4
MRC05         42         M         1.3         492         B         <20         13.0         6.9         6.1           MRC06         50         F         0.9         226         C         <20	MRC03	33	М	0.7	114	F1	<20	30.7	16.0	14.6
MRC06         50         F         0.9         226         C         <20         NA         NA         3.7           MRC07         37         M         1.1         382         B         <20	MRC04	51	М	1.0	171	В	<20	NA	NA	18.4
MRC07         37         M         1.1         382         B         <20         7.5         2.4         5.1           MRC08         50         M         0.5         488         B         <20	MRC05	42	М	1.3	492	В	<20	13.0	6.9	6.1
MRC0850M0.5488B<2016.515.21.3MRC0953F0.8102A1<20	MRC06	50	F	0.9	226	С	<20	NA	NA	3.7
MRC0953F0.8102A1<20NANA17.1MRC1140M1.3350B<20	MRC07	37	М	1.1	382	В	<20	7.5	2.4	5.1
MRC1140M1.3350B<208.41.17.3MRC1231M1.0NAB<20	MRC08	50	М	0.5	488	В	<20	16.5	15.2	1.3
MRC1231M1.0NAB<2011.74.27.5MRC1326M1.3601Recomb B/F1<20	MRC09	53	F	0.8	102	A1	<20	NA	NA	17.1
MRC1326M1.3601Recomb B/F1<20NANA7.2MRC1461M0.9211CRF02_AG<20	MRC11	40	М	1.3	350	В	<20	8.4	1.1	7.3
MRC1461M0.9211CRF02_AG<2011.05.15.9MRC1556M0.698B<20	MRC12	31	М	1.0	NA	В	<20	11.7	4.2	7.5
MRC1556M0.698B<20NANA14.7MRC1932M0.8395B<20	MRC13	26	М	1.3	601	Recomb B/F1	<20	NA	NA	7.2
MRC1932M0.8395B<20NANA1.4MRC2049M0.6294B<20	MRC14	61	М	0.9	211	CRF02_AG	<20	11.0	5.1	5.9
MRC2049M0.6294B<2019.83.416.4MRC2161M0.9179B<20	MRC15	56	М	0.6	98	В	<20	NA	NA	14.7
MRC2161M0.9179B<2030.65.824.8MRC2262M0.8196B<20	MRC19	32	М	0.8	395	В	<20	NA	NA	1.4
MRC2262M0.8196B<2011.63.08.7MRC2358M0.8182B<20	MRC20	49	М	0.6	294	В	<20	19.8	3.4	16.4
MRC2358M0.8182B<2017.00.716.3MRC2454M0.9231B<20	MRC21	61	М	0.9	179	В	<20	30.6	5.8	24.8
MRC2454M0.9231B<2015.62.812.8MRC2548M0.7361B<20	MRC22	62	М	0.8	196	В	<20	11.6	3.0	8.7
MRC25         48         M         0.7         361         B         <20         14.1         2.6         11.5           JZG3034         39         M         0.7         356         B         <20	MRC23	58	М	0.8	182	В	<20	17.0	0.7	16.3
JZG3034 39 M 0.7 356 B <20 3.0 0.5 2.4	MRC24	54	М	0.9	231	В	<20	15.6	2.8	12.8
	MRC25	48	М	0.7	361	В	<20	14.1	2.6	11.5
STAR10 55 M 0.7 327 B <20 21.3 3.0 18.3	UZG3034	39	М	0.7	356	В	<20	3.0	0.5	2.4
	STAR10	55	М	0.7	327	В	<20	21.3	3.0	18.3

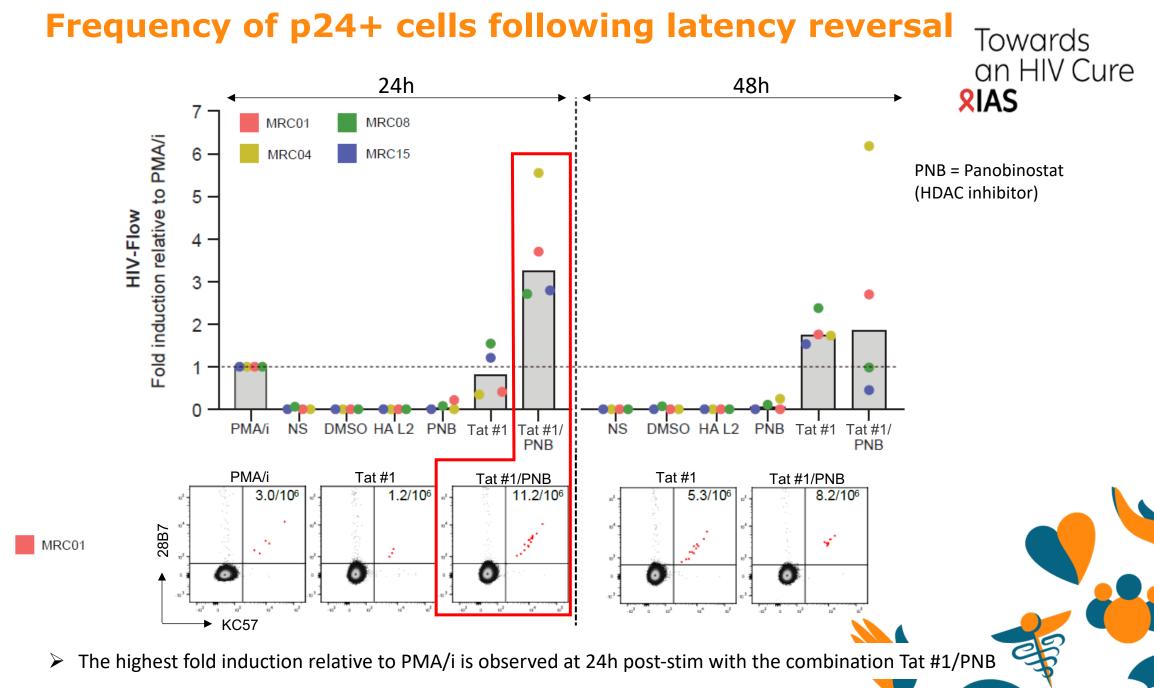


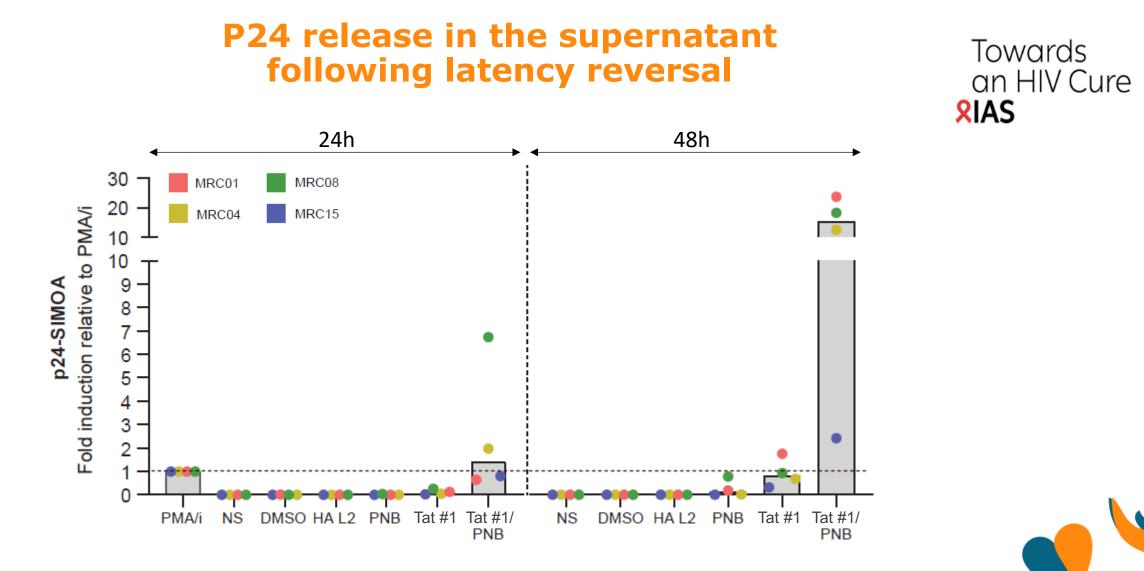
**Viability of the cells** 

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Viability of CD4 T cells is not affected by Tat #1 treatment (both at 24h and 48h)

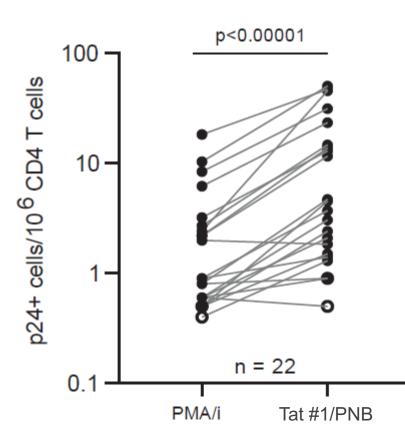
Cell viability is impaired following PNB treatment at 48h





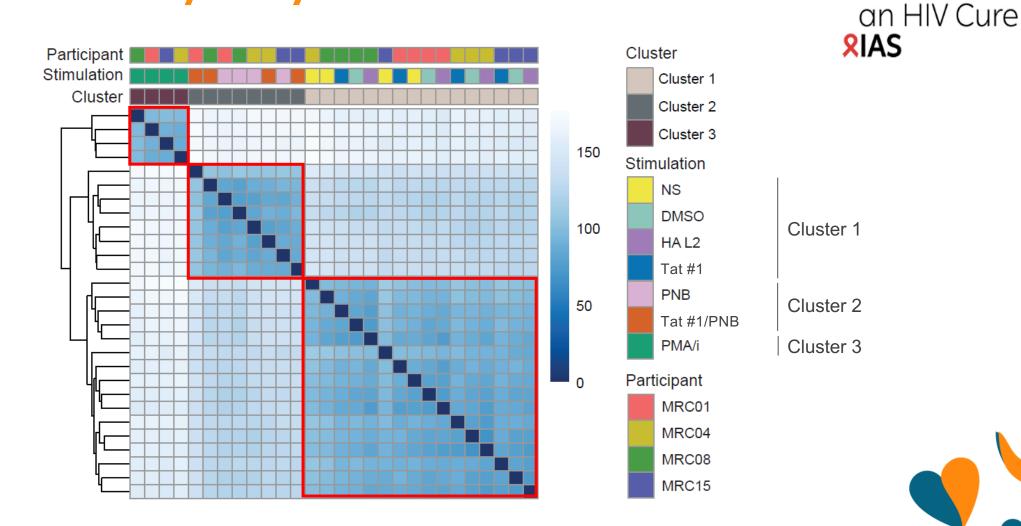
Stimulation with Tat #1 leads to viral particles release in the culture supernatant

#### Frequencies of p24+ cells following stimulation with PMA/i vs Tat #1/PNB Towards an HIV Cure \$IAS



Significantly higher frequencies of p24+ cells are observed following Tat #1/PNB stimulation when compared to PMA/i (median fold increase = 3.9)

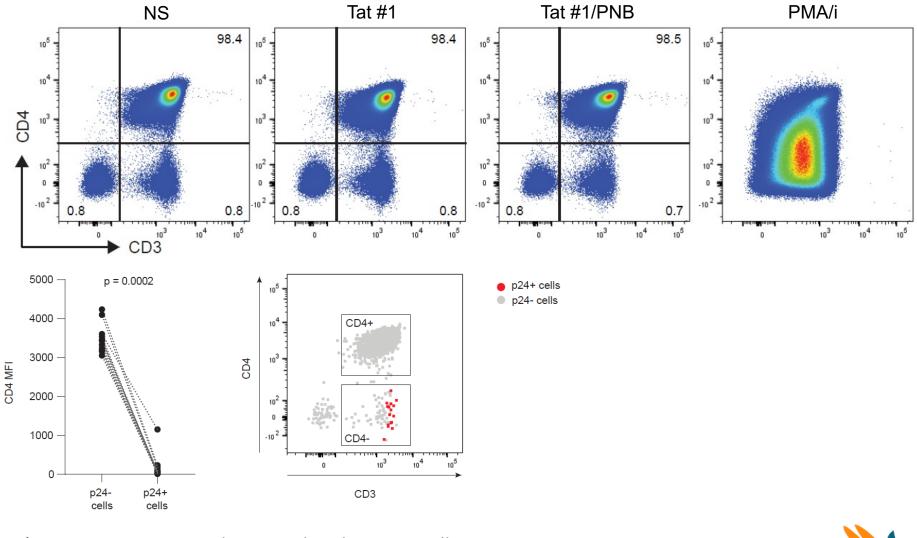
#### **Microarray analyses on bulk CD4 T cells**



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> Tat #1 has a minimal impact on the transcriptomic profile of the cells (clusters with the negative controls)

#### Phenotype of p24+ cells following Tat #1/PNB treatment



CD4 expression is downregulated in p24+ cells

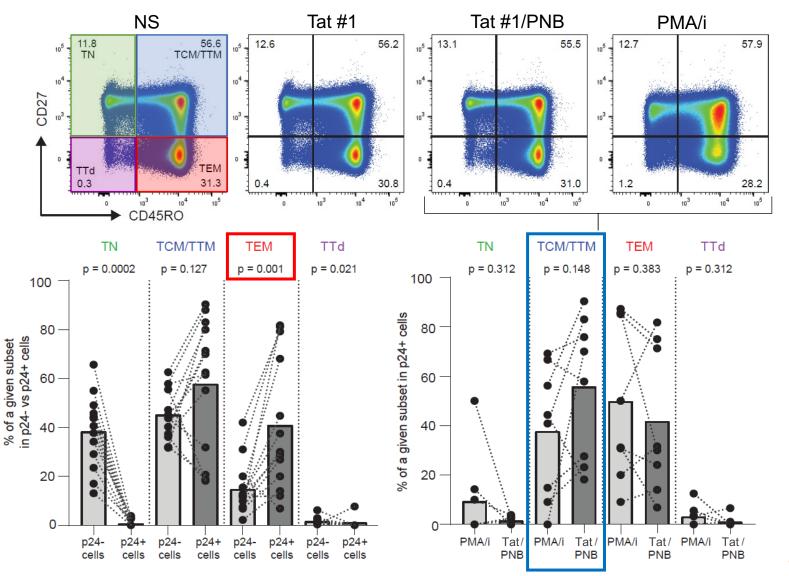


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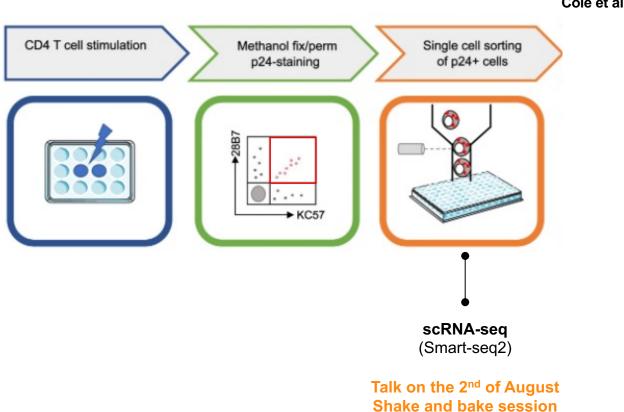
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#### Phenotype of p24+ cells following Tat #1/PNB treatment



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#### Studying the genetic and transcriptional environment Towards of proviruses in p24+ cells an HIV Cure

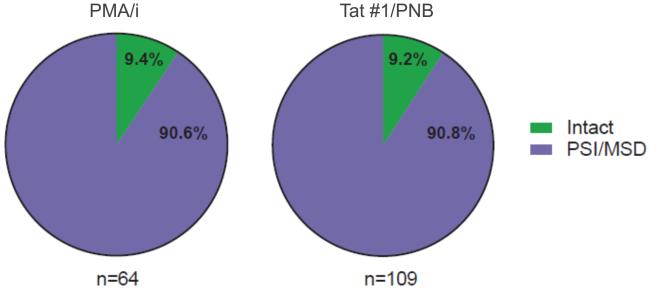


STIP-Seq Cole et al, Nat Com, 2021



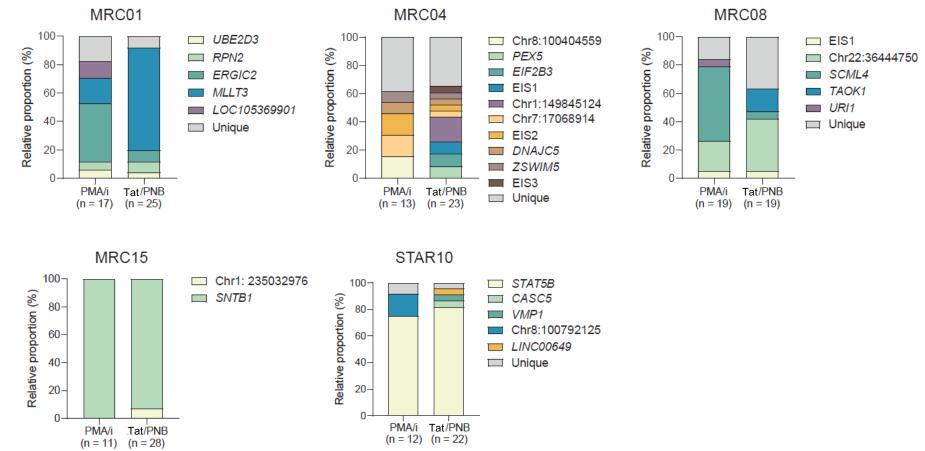
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### Analysis of near-full length sequences in p24+ cells following latency reversal Towards an HIV Cure XIAS



- Most of the p24+ cells harbor PSI/MSD defective proviruses
- Proportions of genome-intact or PSI/MSD-defective proviruses do not differ significantly after PMA/i or Tat #1/PNB stimulation

#### Analysis of the integration sites in p24+ cells following latency reversal



- Tat #1/PNB globally reactivates the same clones as PMA/i
- Some rare and minor clones were detected only in one of the two conditions
- Some clones are represented in different proportions between the two conditions (e.g. MLLT3, SCML4)



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# Conclusions



- Tat : physiologically relevant, not toxic *in vitro*
- Does not modify the transcriptome/phenotype of the cells
- In combination with PNB:
  - It induces latency reversal in a higher proportion of latently infected cells than PMA/i
  - Reactivates the same clones than PMA/i, with some exceptions
- Can be used as a tool to study:
  - The proviral sequence and integration site in p24+ cells
  - Phenotype of p24+ cells

## **Acknowledgements**

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Liège university

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Flow cytometry and sequencing cores from Ghent and Janssen



