

Delivery is key



- **The problem**: Delivery to target tissues is the primary obstacle impeding the clinical translation of OligoNucleotide (ON) therapeutics
- The solution? Cell Penetrating Peptides (CPPs) are increasingly prominent in the quest to address this delivery obstacle
- They've been around for 40 years with only incremental progress demonstrated what's different this time?
 - Unimproved 'hit' sequences that are more than 10x as efficient as 'conventional CPPs'
 - Evidence of tissue tropism to target specific cell types concentrating drug in diseased tissues
 - Developments in analytics frameworks that enable multi-variate optimisation (improving 'hit to lead' and development conversion)
 - Early transition to 'build the delivery platform' in the context of the cargo essential in the context of cargo-specific properties
 - Indications that absolute disease correction thresholds are within reach (eg. Sarepta's pipeline of PPMO therapies)

A tale of two companies...



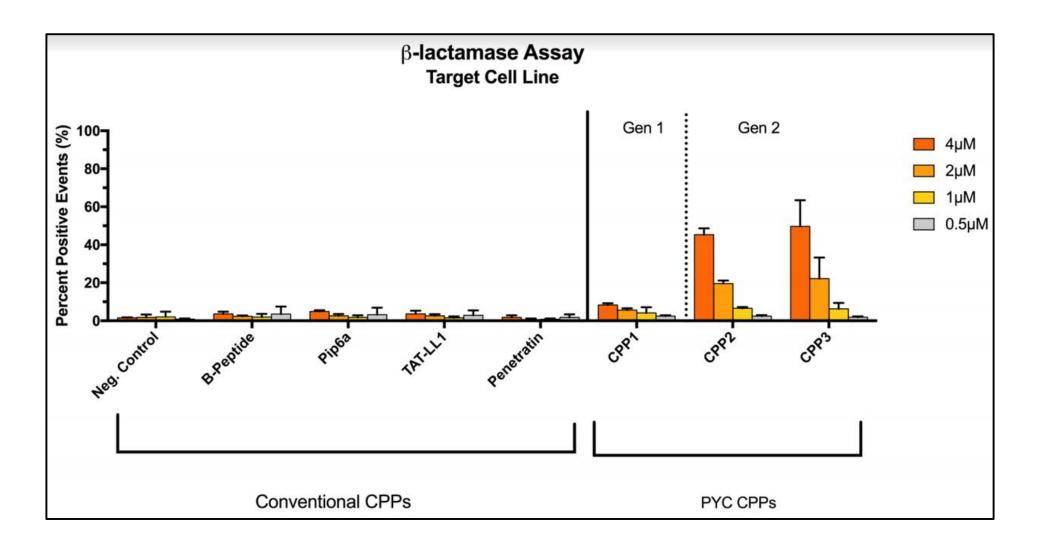
Company	Parental CPP sequences	CPP sequences evaluated in lead selection	Development stage
SAREPTA THERAPEUTICS	1*	4	 1 clinical PPMO 5 IND submissions for PPMOs planned in 2019
PHYL <mark>Ŏ</mark> GICA	1,000 – 10,000	~100	Pre-clinical – therapeutic <i>in vivo</i> validation
We are balancing the need to rapidly develop the platform with			

the value of carefully considering the breadth of data we have

^{*} Sarepta used the B-peptide (a modified poly-arginine CPP) as their starting sequence

Phylogica's CPPs are more efficient than 'conventional CPPs'



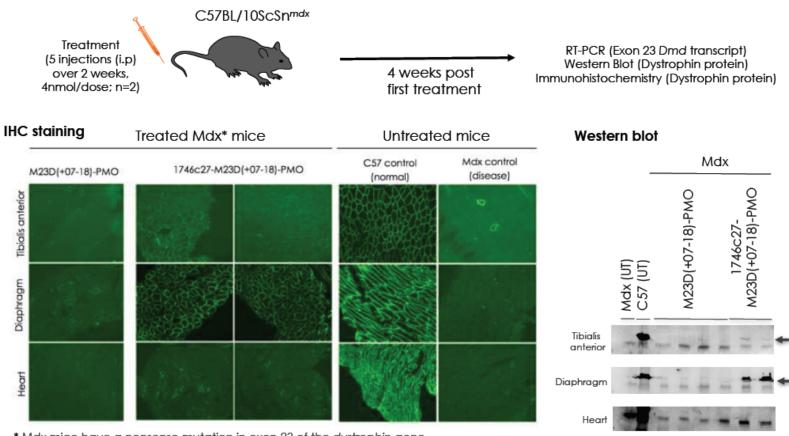


Our '1st generation' CPPs are effective in vivo - '2nd gen' results will follow soon



Dystrophin levels are restored by CPP-mediated delivery of DMD PMO in vivo

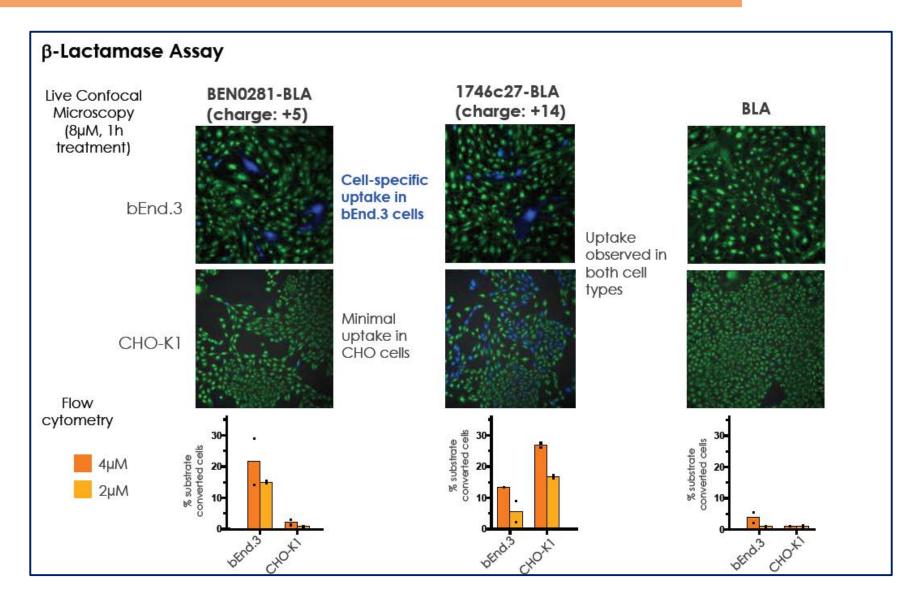
Sustained response in CPP-PMO treated mice



^{*} Mdx mice have a nonsense mutation in exon 23 of the dystrophin gene

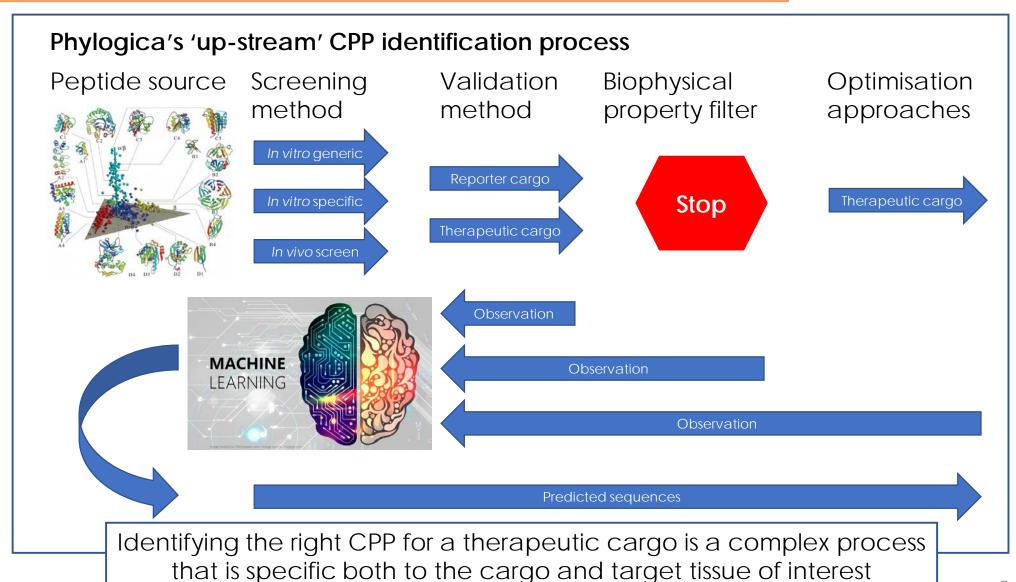
Indications of cell-specific CPPs suggest further concentration of cargo is possible





Recent developments in analytics allow us to consider many more variables





Is the juice worth the squeeze?



- The richness of the data set that we generate from the Phylomer libraries is both a blessing and a curse:
 - A blessing in the number and quality of CPPs that we identify (Phylogica's CPPs substantially outperform the 'conventional CPP landscape'); and
 - A curse in the amount of data that we are handling screening and validating
 CPPs is both expensive and time-consuming
- We have the privileged ability to see the 'macro rule' (the link between the properties
 of a CPP and its ability to deliver a cargo) across CPPs of divergent characteristics
 (raising the possibility that we can combine 'parts' of CPPs to create high performing
 'chimeras')
- We can also define the 'micro' rule through creation of variants of parental sequences to 'fine tune' a CPP
- BUT...
- These activities must be conducted in the context of the cargo of interest (and ideally the target cell of interest too) - we are building a platform tailored for ONs

In summary...



- CPPs have the ability to open up new target tissues/cells and disease indications for ON therapeutics
- Phylogica's platform can deliver a step change in the amount of drug delivered into a target cell by:
 - Using substantially more efficient 'second generation' peptides;
 - Exploiting observed patterns of differential cellular uptake to target diseased cells; and
 - Using advanced analytics to select for drug-like properties
- We are building our platform specifically for ON therapeutics and will be moving into therapeutic in vivo evaluation very shortly



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