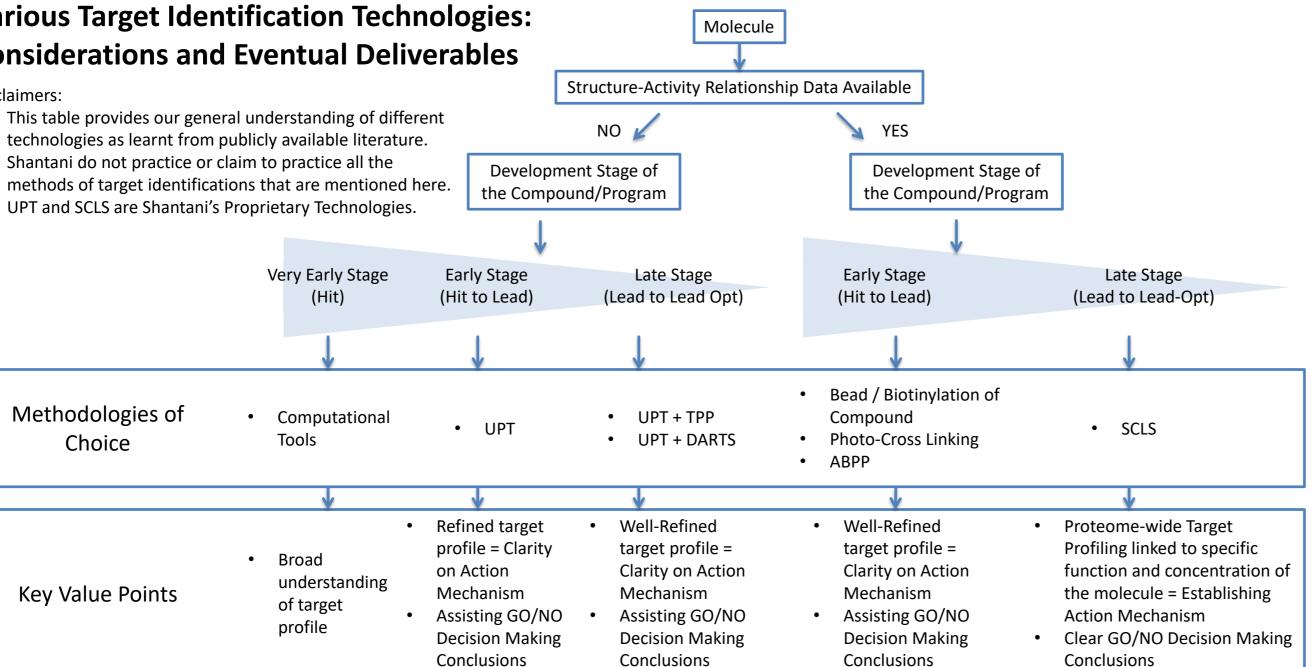
Various Target Identification Technologies: Considerations and Eventual Deliverables

Disclaimers:

- technologies as learnt from publicly available literature.
- methods of target identifications that are mentioned here.



- Conclusions
- Conclusions
- Conclusions

Abbreviation

- **UPT** = Unique Polymer Technology (Shantani's Proprietary Technology)
- **TPP** = Thermal Proteome Profiling Savitski et al., Science (2014) 346 (6205):1225784)
- **DARTS** = Drug Affinity Responsive Target Stability (Lomenick et al. PNAS (2009) 106:21984-9)
- **ABPP** = Activity Based Protein Profiling (Evans et al. Nat Biotechnol (2005) 23:1303-7)
- **SCLS** = Subcellular Location Specific Target Capture & Identification Technology (Shantani's Proprietary Technology)

For Other Available Target Identification Methodologies:

Saxena et al. Expert Opin. Drug Discov.

(2009) 4:701-714

Saxena C. Expert Opin. Drug Discov.

(2016) 11:1017-1025

Target Identification: Comparison of Chemical-Proteomics Based Mass-Spectrometry Based Methodologies

	Resources	es Molecule Derivatization Not Needed / Structure-Activity Relationship not Available				Molecule Needs to be Derivatized / Structure-Activity Relationship Available			
Criteria		Computational Tools	UPT	ТРР	DARTS	Bead / Biotinylation	Photo-Cross Linking	АВРР	SCLS
Resources	Chemistry	-	-	-	-	++	+++	+++	++
	Computational and/OR Statistical	+++++	++	++++	++++	++	+++	+++	++
	Biochemistry	-	+	+	+	+	+	+	++
	Cell-Biology	-	+	+	+	+	+	+	++
	Mass- Spectrometry	-	++	+++	+++	+++	+++	+++	++
Time		+	++	++	++	++++	++++	++++	+++++
Cost		+	+++	++++	++++	+++	++++	++++	+++++
False Positive Identification		~ 80 %	~40%	~60%	~60%	~40%	~40%	~30%	~20%
Key Advantage(s)		Quick & Cost Effective	Underivatized bait- molecule Target Identification through affinity based target enrichment	Underivatized bait- molecule	Underivatized bait- molecule	Targets of varied affinity (high and low) are captured	Covalent capture of target	Activity based target capture	Capture of target from live cell in sub-cellular location specific manner
Disadvantages		Very High False Positive Rate	Certain highly water Soluble Compounds can not used	 Limited to Soluble Proteins only Complex data- analysis 	Limited to Highly Abundant Proteins only	 Bait-molecule derivatization is needed for immobilization Limited success in capturing membrane targets 	 Bait-molecule derivatization is needed for immobilization Primary bait-molecule may loose activity because relatively larger size of probes 	 Bait-molecule derivatization is needed for immobilization Very Specific probes needs to be synthesized –target class specific capture 	 Bait-molecule derivatization is needed for immobilization Particular Function / phenotype linked target are captured