

Design and discovery of novel Granzyme B ligands and evaluation as diagnostic tools in CAR T-cell therapy assessment

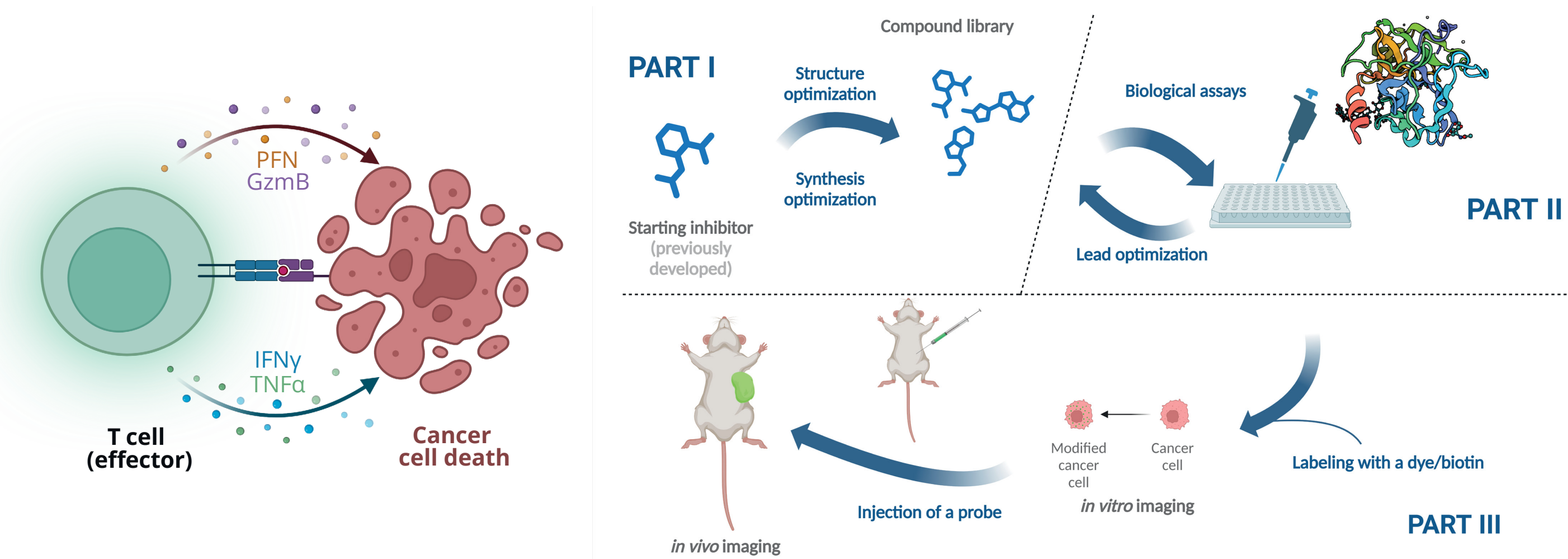


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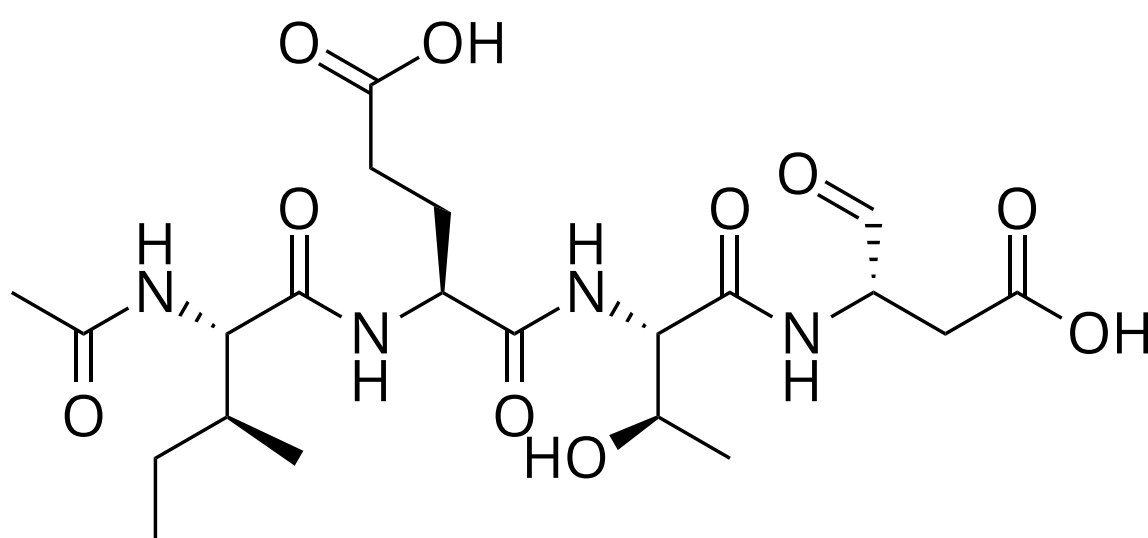
INTRODUCTION

- Granzyme B (GzmB) is a serine protease expressed in cytotoxic T-cells and natural killer (NK) cells that targets virus-infected cells and tumor cells.¹
- GzmB activates the apoptotic pathway by cleavage of caspases 8, 10, 3, 7.^{1,2}
- GzmB acts as a signal factor in CAR T-cell immunotherapy.
- GzmB represents an intra-tumoral and extra-tumoral target for molecular imaging for early cancer diagnostics and immunotherapy efficiency.³



AIMS AND OBJECTIVES

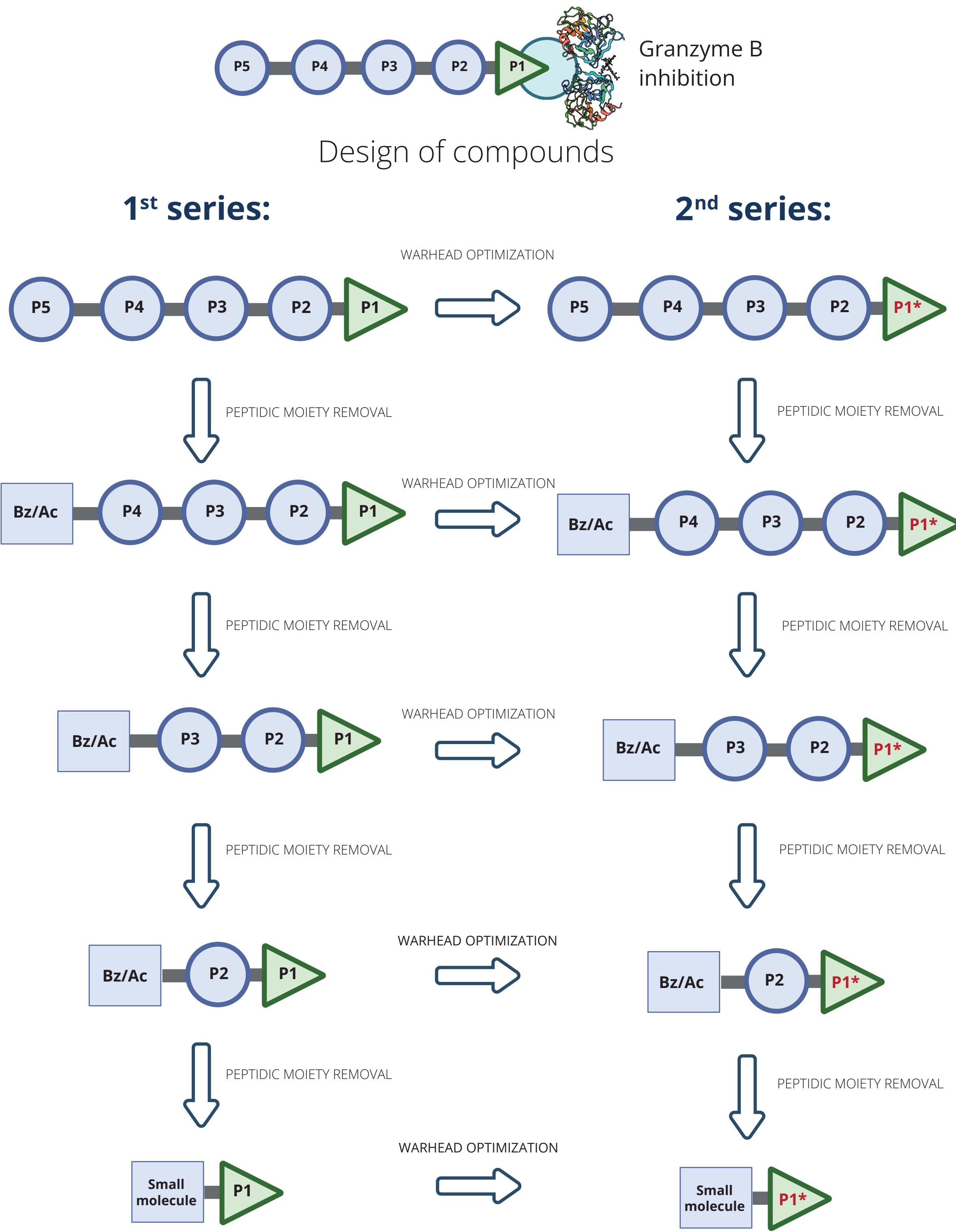
- Synthesis of a library of novel GzmB inhibitors with a more drug-like profile to overcome metabolic stability issue.
- In vitro* biological evaluation of the compounds against human GzmB, caspase-3, cathepsins B, L, and S.
- Generation of structure-activity relationship (SAR) data.
- Selection of the most suitable (most potent & selective) inhibitor to be converted into a probe (attachment of a linker with a tag).



IEDT inhibitor³
 $K_i = 80 \text{ nM}$

METHOD AND RESULTS

1. COMPOUNDS LIBRARY SYNTHESIS



- 50+ novel inhibitors featuring P1-P5 modifications and diverse linker design are synthesized and characterized.



2. BIOLOGICAL ASSESSMENT

Granzyme B expression: (performed by dr. Emile Verhulst & dr. Fernanda Rocho)

Source: HEK293 T-cells (transient transfection).

Activity tests:

Substrate: Ac-IEPD-pNA at the K_M value.

Enzyme: recombinant human Granzyme B.

~ 10 hit compounds observed ($IC_{50} < 500 \text{ nM}$).

Reference molecules

Compound	IC_{50} (nM)	Note
M1	>2245	Main structural motif
M2	125 ± 5	Ref. molecule 1
M3	46 ± 3	Ref. molecule 2

Polarity changes

Compound	IC_{50} (nM)	Note
M4	36 ± 7	Polar
M5	280 ± 3	Non-polar

Linker series

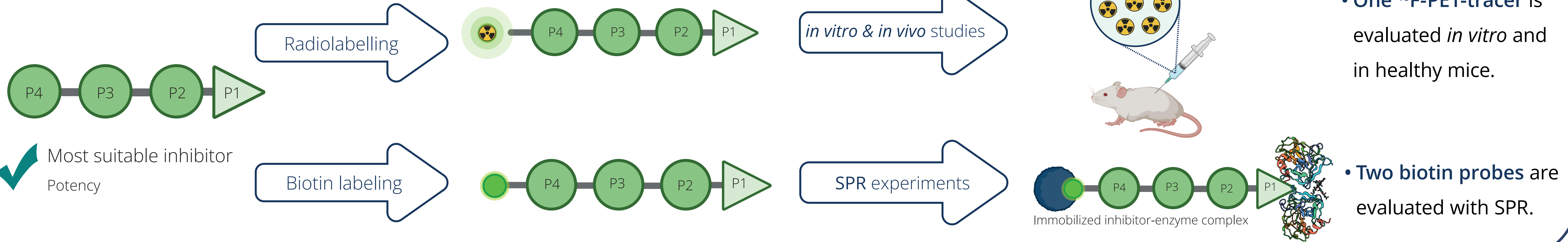
Compound	IC_{50} (nM)	Note
M6	452 ± 4	Linker 1
M7	51 ± 3	Linker 1 ↑ polarity
M8	55 ± 1	Linker 2 ↑ polarity
M9	28 ± 3	Linker 3 ↑ polarity
M10	160 ± 33	Linker 4
M11	93 ± 15	Linker 4 - cold PET-probe
M12	187 ± 14	Linker 2 - Biotin probe

Selectivity tests:

- High selectivity ($SI \gg 100$) against Cathepsins B, L and S.
- Moderate selectivity ($SI \approx 10$) over Caspase-3.

- Compounds library screened against Granzyme B, Caspase-3, and Cat B/L/S.

3. PROBES DEVELOPMENT



REFERENCES

- Angew. Chem. Int. Ed. 2021. 60 (11), 5699–5703.
- J. Biol. Chem. 2020. 295 (28), 9567–9582.
- Nature communications. 2021. 12(1), 1-14.
- Experimental & molecular medicine. 2018. 50(5), 1-11.



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