

We claim:

1. A method of identifying a ligand, comprising:
 - (a) contacting a biological molecule and a probe molecule; wherein the probe molecule comprises a binding element, a reporter group, and a reactive moiety; the probe molecule binds to the biological molecule via the binding element; and the reactive moiety forms a covalent bond with the biological molecule, thereby forming a conjugate;
 - (b) contacting the conjugate and a detectable molecule comprising a functional moiety that reacts with the reporter group, thereby forming a detectable conjugate;
 - (c) contacting the detectable conjugate and a solid support; wherein the solid support comprises a recognition moiety; and the recognition moiety binds to the detectable conjugate, thereby forming a bound detectable conjugate; and
 - (d) detecting the bound detectable conjugate, thereby identifying the probe molecule as a ligand for the biological molecule.
2. The method of claim 1, wherein the binding element is a small molecule.
3. The method of claim 1, wherein the binding element is a peptide.
4. The method of claim 1, wherein the binding element is a nucleic acid.
5. The method of claim 4, wherein the nucleic acid is RNA or DNA.
6. The method of any one of claims 1-5, wherein the binding element is a component of a library comprising a plurality of binding elements.
7. The method of claim 6, wherein the binding elements are small-molecule fragments; and each small-molecule fragment has at least one of the following characteristics: molecular weight ≤ 300 Da, cLogP ≤ 3 , hydrogen bond donors ≤ 3 , and hydrogen bond acceptors ≤ 3 .
8. The method of any one of claims 1-7, wherein the biological molecule is a protein.
9. The method of claim 8, wherein the reactive moiety forms a covalent bond with an amino acid of the protein.

10. The method of any one of claims 1-7, wherein the biological molecule is a lipid.
11. The method of any one of claims 1-7, wherein the biological molecule is a carbohydrate.
12. The method of any one of claims 1-7, wherein the biological molecule is a nucleic acid.
13. The method of claim 12, wherein the nucleic acid is RNA.
14. The method of claim 12, wherein the nucleic acid is DNA.
15. The method of any one of claims 1-14, wherein the biological molecule comprises an epitope tag.
16. The method of claim 15, wherein the epitope tag is FLAG, 6xHis, HA, c-myc, glutathione-S-transferase, Strep-tag, maltose-binding protein, chitin-binding protein, S-tag, V5 tag, or AviTag.
17. The method of any one of claims 1-16, wherein the reporter group comprises an azadibenzocyclooctyne, a thiol, an alkene, an alkyne, an azide, a tetrazine, a *trans*-cyclooctene, a (diphenylphosphino)aryl, (diphenylphosphino)alkyl, or an activated ester (e.g., a hydroxybenzotriazole (HOBt) ester).
18. The method of any one of claims 1-17, wherein the reactive moiety is a photocrosslinker group, a sulfonyl fluoride, a fluorosulfate, a Michael acceptor moiety, a leaving group moiety, or a moiety that forms a covalent bond with a nucleophilic moiety in the side chain of a naturally occurring alpha amino acid (e.g., with the thiol group of a cysteine, the amino group of a lysine, the hydroxyl group of a serine or threonine, or the phenol group of a tyrosine).
19. The method of any one of claims 1-18, wherein the detectable molecule comprises digoxigenin, nickel NTA (nitrilotriacetic acid), a chromophore, or a luminophore.
20. The method of claim 19, wherein the detectable molecule comprises a digoxigenin.

21. The method of claim 19, wherein the detectable molecule comprises a nickel NTA (nitrilotriacetic acid).
22. The method of claim 19, wherein the detectable molecule comprises a chromophore.
23. The method of claim 22, wherein the chromophore comprises non-fluorochrome chromophore, quencher, an absorption chromophore, fluorophore, organic dye, inorganic dye, metal chelate, or a fluorescent enzyme substrate.
24. The method of claim 19, wherein the detectable molecule comprises a luminophore.
25. The method of any one of claims 1-18, wherein the detectable molecule is biotin.
26. The method of claim 25, wherein the biotin is bound to a streptavidin conjugate.
27. The method of claim 26, wherein the streptavidin conjugate comprises a HRP, a SulfoTag, a fluorophore, or a metal chelate.
28. The method of any one of claims 1-27, wherein the functional moiety comprises an azadibenzocyclooctyne, a thiol, an alkene, an alkyne, an azide, a tetrazine, a *trans*-cyclooctene, a (diphenylphosphino)aryl, (diphenylphosphino)alkyl, or an activated ester (e.g., a hydroxybenzotriazole (HOBt) ester).
29. The method of any one of claims 1-28, wherein the solid support is a membrane, glass, plastic, synthetically prepared polymer, an eppendorf tube, a well of a multi-well plate, or a surface plasmon resonance chip.
30. The method of any one of claims 1-29, wherein the recognition moiety is an antibody, a DNA binding protein, a RNA binding protein, a carbohydrate binding protein, or a lipid binding protein.
31. The method of claim 30, wherein the recognition moiety is an antibody.

32. The method of claim 31, wherein the antibody is an antibody against the biological molecule.
33. The method of claim 31, wherein the antibody is an antibody against the epitope tag.
34. The method of any one of claims 1-33, wherein step (d) comprises detecting the bound detectable conjugates via ELISA.
35. The method of any one of claims 1-33, wherein step (d) comprises detecting the bound detectable conjugates via Western blot.
36. The method of any one of claims 1-33, wherein step (d) comprises detecting the bound detectable conjugates via immunofluorescence assay.
37. The method of any one of claims 1-33, wherein step (d) comprises detecting the bound detectable conjugates via fluorometric assay.
38. The method of any one of claims 1-33, wherein step (d) comprises detecting the bound detectable conjugates via fluorometric microvolume assay technology (FMAT).
39. The method of any one of claims 1-33, wherein step (d) comprises detecting the bound detectable conjugates via cell subcellular staining.
40. The method of any one of claims 1-39, wherein the method is performed on a crude cellular extract comprising the biological molecule.
41. The method of any one of claims 1-39, wherein the method is performed on a liposomal preparation of proteins comprising the biological molecule.
42. The method of any one of claims 1-39, wherein the method is performed on an isolated organelle comprising the biological molecule.
43. The method of any one of claims 1-39, wherein the method is performed on a purified protein preparation comprising the biological molecule.
44. The method of any one of claims 1-39, wherein the method is performed *in situ*.

45. The method of any one of claims 1-39, wherein the method is a cell-based assay.
46. The method of claim 45, wherein the biological molecule is expressed in a cell.
47. The method of claim 46, wherein the cell is engineered to express the biological molecule.
48. The method of any one of claims 45-47, wherein the cell is lysed prior to step (b).
49. The method of any one of claims 1-48, wherein step (d) further comprises quantifying the amount of the bound detectable conjugate.
50. The method of claim 49, wherein step (a) further comprises a substrate for the biological molecule.
51. The method of claim 50, wherein the amount of the bound detectable conjugate formed in the presence of the substrate for the biological molecule is less than the amount of the bound detectable conjugate formed in the absence of the substrate (i.e., the probe molecule is a substrate-competitive probe).
52. The method of claim 50, wherein the amount of the bound detectable conjugate formed in the presence of the substrate for the biological molecule is greater than the amount of the bound detectable conjugate formed in the absence of the substrate (i.e., the probe molecule is a substrate-cooperative probe).