WHAT IS CLAIMED IS:

1. A system comprising:
   (a) an array comprising a plurality of capture probes, wherein a capture probe of the plurality of capture probes comprises: (i) a spatial barcode and (ii) a capture domain;
   (b) one or more semi-porous materials disposed between a biological sample and the array; and
   (c) an electric field, wherein the electric field promotes the migration of one or more analytes in the direction of the array, wherein the capture domain of the capture probe binds to a first analyte of the one or more analytes, and wherein the first analyte is a nucleic acid.

2. The system of claim 1, wherein a first semi-porous material of the one or more semi-porous materials retains one or more second analytes.

3. The system of claim 1 or 2, wherein the first semi-porous material comprises a plurality of analyte-binding moieties, wherein an analyte-binding moiety of the plurality specifically binds to a second analyte of the one or more second analytes and wherein the analyte binding moiety is an antibody or antigen-binding fragment thereof.

4. The system of claim 2 or 3, wherein the first analyte is mRNA and the second analyte is a protein.

5. The system of any one of claims 1-4, further comprising determining the location of the second analyte in the first semi-porous material, wherein determining the location of the second analyte in the first semi-porous material comprises immunofluorescence staining.

6. The system of claim 5, wherein determining the location of the second analyte in the first semi-porous material comprises:
   obtaining an image of the first semi-porous material; and

correlating the immunofluorescence staining in the image of the first semi-porous material with an image of the biological sample.

7. The system of any one of claims 1-6, wherein the one or more semi-porous materials comprises a second semi-porous material, wherein the second semi-porous material retains one or more third analytes.

8. The system of claim 7, wherein the second semi-porous material comprises a plurality of analyte-binding moieties, wherein an analyte binding moiety of the plurality of analyte binding moieties specifically binds to a third analyte of the one or more third analytes.

9. The system of claim 8, wherein the third analyte comprises a second nucleic acid species.

10. The system of claim 8 or 9, further comprising determining the location of the third analyte in the second semi-porous material, wherein determining the location of the third analyte in the second semi-porous material comprises the use of *in situ* hybridization.

11. The system of claim 10, wherein determining the location of the third analyte comprises:
obtaining an image of the second semi-porous material; and
correlating the *in situ* hybridization in the image of the second semi-porous material with an image of the biological sample.

12. The system of claim 8, further comprising crosslinking the one or more third analytes to the second semi-porous material, and optionally, cross-linking one or more second analytes to the first porous material.

13. The system of any one of claims 8-12, further comprising removing the second semi-porous material after the second semi-porous material retains the one or more third analytes.

14. The system of any one of claims 1-6, further comprising:

removing the first semi-porous material after the first semi-porous material retains the one or more second analytes; and

disposing the second semi-porous material between the biological sample and the array.

15. The system of claim 14, wherein the electric field is applied after removal of the first semi-porous material and disposal of the second semi-porous material between the biological sample and the array and wherein the second semi-porous material retains one or more third analytes.

16. The system of any one of claims 1-15, wherein the capture probe further comprises one or more of: a cleavage domain, a functional domain, and a unique molecular identifier.

17. The system of any of claims 1-16, further comprising imaging the biological sample.

18. The system of any one of claims 1-17, wherein the semi-porous material comprises a hydrogel or a permeable membrane.

19. The system of any one of claims 1-18, wherein the one or more semi-porous materials have non-uniform pore sizes.

20. The system of any one of claims 1-18, wherein the one or more semi-porous materials have a substantially uniform pore size.

21. The system of any one of claims 1-20, wherein the one or more semi-porous materials comprise one or more fiducial markers that align the one or more semi-porous materials to the image of the biological sample.

22. The system of any one of claims 1-21, wherein the biological sample is a tissue section or a fixed tissue section.

23. The system of any one of claims 1-22, further comprising a step of fixing and/or permeabilizing the biological sample.

24. The system of any one of claims 1-23, wherein an electric field is applied to a discrete area of the biological sample or a discrete area of the array.

25. The system of any one of claims 1-24, wherein, the biological sample, the one or more semi-porous materials, and/or the array, are in direct contact with a buffer.

26. The system of any one of claims 1-25, wherein the system further comprises determining (i) the sequence of the spatial barcode, or a complement thereof, and (ii) all or a portion of the sequence of the first analyte, or a complement thereof, and using the sequences of (i) and (ii) to determine the location of the first analyte in the biological sample.

27. A system comprising:

(a) a first array comprising a plurality of first capture probes, wherein a first capture probe of the plurality of first capture probes comprises: (i) a first spatial barcode and (ii) a first capture domain;

(b) a semi-porous material, wherein the semi-porous material is disposed between the biological sample and the first array;

(c) a first electric field, wherein the first electric field promotes the migration of one or more analytes in the direction of the first array, wherein the first capture domain specifically binds to a first analyte of the one or more analytes, and wherein the semi-porous material retains a second analyte of the one or more analytes;

(d) a second array comprising a plurality of second capture probes, wherein a second capture probe of the plurality of second capture probes comprises: (i) a second spatial barcode and (ii) a second capture domain;

(e) a second electric field, wherein the second electric field promotes the migration of the second analyte in the direction of the second array, wherein the second capture domain specifically binds to an analyte capture sequence; and

(f) determining (i) the sequence of the second spatial barcode, or a complement thereof, and (ii) all or a portion of the sequence of the second analyte, or a complement thereof, and using the sequences of (i) and (ii) to determine the location of the second analyte in the biological sample.

28. The system of claim 27, wherein the capture probe further comprises one or more of: a cleavage domain, a functional domain, and a unique molecular identifier.

29. The system of claim 27 or 28, wherein the system further comprises imaging the biological sample.

30. The system of any one of claims 27-29, wherein the semi-porous material comprises a hydrogel or a permeable membrane.

31. The system of any one of claims 27-30, wherein the semi-porous material has non-uniform pore sizes.

32. The system of any one of claims 27-30, wherein the semi-porous material has a substantially uniform pore size.

33. The system of any one of claims 27-32, wherein the semi-porous material comprises one or more fiducial markers that align the semi-porous material to the image of the biological sample.

34. The system of any one of claims 27-33, wherein the biological sample is a tissue section or a fixed tissue section.

35. The system of any one of claims 27-34, further comprising a step of fixing and/or permeabilizing the biological sample.

36. The system of any one of claims 27-35, wherein the first electric field, the second electric field, or both, is applied to a discrete area of the biological sample.

37. The system of any one of claims 24-36, wherein the first electric field is applied to a discrete area of the first array and wherein the second electric field is applied to a discrete area of the second array.

38. The system of any one of claims 27-37, wherein the biological sample, the semi-porous material, and/or the first array are in direct contact with a buffer.

39. The system of any one of claims 27-37, wherein the biological sample, the semi-porous material, and/or the second array are in direct contact with a buffer.

40. The system of any one of claims 27-39, wherein the semi-porous material separates one or more analytes from other molecules in the biological sample.

41. The system of any one of claims 27-40, wherein the semi-porous material comprises one or more analyte binding moieties, wherein the one or more analyte binding moieties are crosslinked to the semi-porous material and bind a nucleic acid.

42. The system of any one of claims 27-41, wherein the semi-porous material comprises one or more analyte capture agents, wherein the one or more analyte capture agents are crosslinked to the semi-porous material and wherein an analyte capture agent comprises:

(i) an analyte binding moiety, wherein the analyte binding moiety comprises an antibody or an antigen-binding fragment thereof,

(ii) an analyte binding moiety barcode, wherein the analyte binding moiety barcode identifies the analyte binding moiety; and

(iii) the analyte capture sequence, wherein the analyte capture sequence binds to the second capture domain of the second capture probe.

43. The system of claim 42, wherein the first analyte is a mRNA and the second analyte is a protein.

44. The system of any one of claims 27-43, wherein the system further comprises:

(a) determining (i) the sequence of the first spatial barcode, or a complement thereof, and (ii) all or a portion of the sequence of the first analyte, or a complement thereof, and using the sequences of (i) and (ii) to determine the location of the first analyte in the biological sample; and

(b) determining (i) the sequence of the second spatial barcode, or a complement thereof, and (ii) all or a portion of the sequence of the analyte binding moiety barcode, or a complement thereof, and using the sequences of (i) and (ii) to determine the location of the second analyte in the biological sample.