

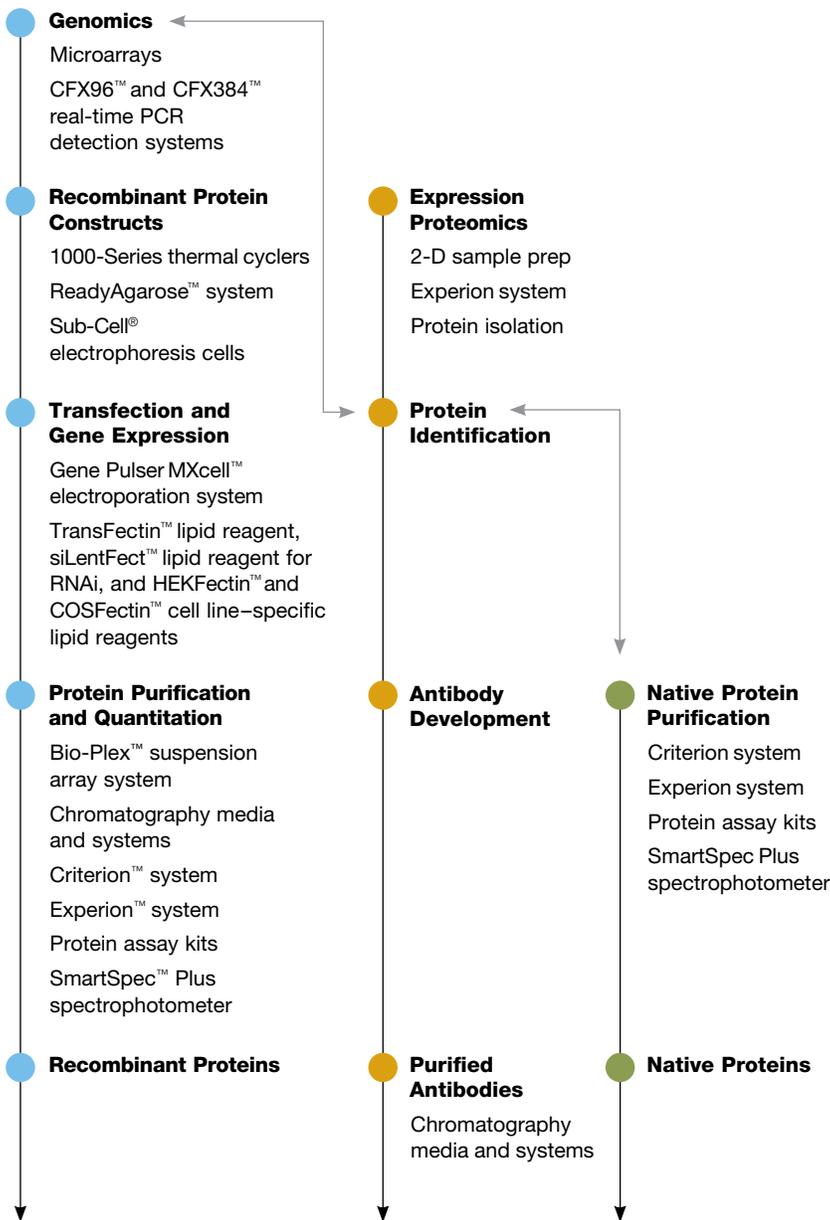
ProteOn™ XPR36
Protein Interaction Array System



The Power of Parallel Processing



Bridge the Gap Between Discovery and Function



Performance

- XPR™ technology — a crisscross 6 x 6 microfluidic array for analysis of up to 36 biomolecular interactions
- Parallel processing for efficient optimization of immobilization and interaction conditions
- Temperature control of autosampler and sensor chip

Sensor Chips

- Protein-protein, protein-peptide, protein-small molecule, and protein-DNA interaction analysis
- Easy activation for efficient immobilization and high ligand activity
- Uniform spot-to-spot response
- Bar codes for automatic chip identification and usage record

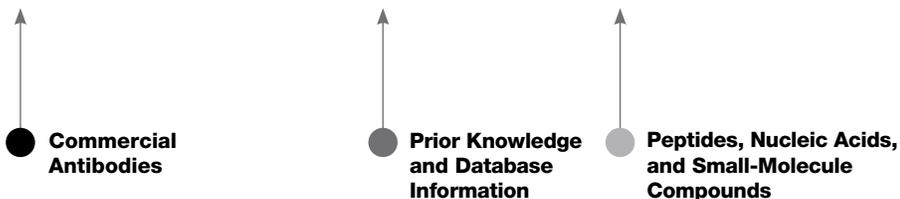
Software

- Flexible, user-guided, wizard-driven software and maintenance protocols
- Protocol templates to assist with experimental design and setup
- Interspot referencing and reference channel subtraction
- Automatic baseline and injection alignment of sensorgrams
- 3 analysis modules: determination of kinetic constants, concentration determination, and equilibrium analysis

Regulatory Tools

- ProteOn Manager, Security Edition, for U.S. FDA 21 CFR Part 11 compliance
- ProteOn XPR36 Installation Qualification/Operation Qualification (IQ/OQ) Kit

▪ Concentration ▪ Equilibrium ▪ Kinetics ▪ Specificity/Affinity





Proteins play a crucial role in cellular structure and function. The search, discovery, and characterization of proteins is the first step in understanding how each protein participates in a biological process. The next logical step is to understand how a protein functions: What binding partners exist? Is this protein involved in a network? How does one protein interface with another? What are the binding dynamics? The ProteOn XPR36 system bridges the gap between discovery and function, providing the ability to understand protein interactions with the efficiency, flexibility, and versatility of parallel processing.

ProteOn XPR36 Protein Interaction Array System



The ProteOn XPR36 protein interaction array system is an SPR optical biosensor designed to provide all the benefits of parallel processing. XPR technology — a dynamic and unique approach to multiplexing — greatly improves the efficiency, flexibility, and results of your experimental design, enabling you to run more experiments in a shorter period of time. The ProteOn XPR36 workflow is guided by ProteOn Manager™ software — an easy-to-use, intuitive interface that provides a flexible yet guided approach to instrument control, experimental setup, and data analysis. The ProteOn XPR36 system includes all the components necessary for successful and efficient protein interaction analysis: instrumentation, software, sensor chips, buffers and reagents, protocol development kits, and tools to meet regulatory requirements.



One-shot Kinetics™

Achieve a complete kinetic profile of a biomolecular interaction:

- In a single experiment
- On a single chip
- Without the need for regeneration

Optimize, Maximize, Analyze

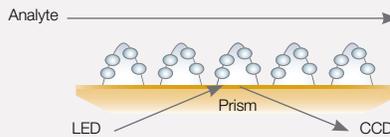
XPR technology offers the ability to monitor up to 36 different interactions in real time by processing, in parallel, 6 different ligands with 6 different analytes.

The 6 x 6 interaction array provides:

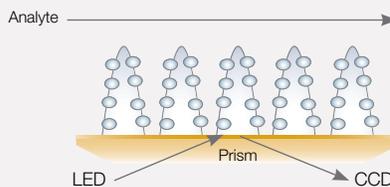
- Flexible experimental design — test a variety of experimental conditions on a single chip in a single experiment
- High sample throughput — immobilize a single panel of 6 ligands and screen multiple panels of analytes
- Rapid data generation — screen up to 180 interactions per hour with immobilization of 6 ligands

1. Select a Sensor Chip Surface

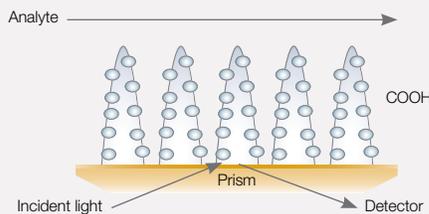
A family of surface chemistries is available for easy ligand activation. ProteOn sensor chips have been thoroughly tested for a wide range of protein-protein, protein-peptide, protein-small molecule, and protein-DNA interactions. Each sensor chip includes a unique bar code for identification and a usage record. ProteOn sensor chips offer outstanding kinetic response characteristics, high binding capacities, sufficient sensitivity to detect low molecular weight analytes, uniform spot-to-spot response, minimal drift, and long-term storage stability.



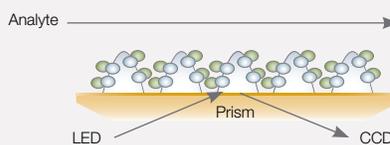
ProteOn GLC sensor chip. For general amine coupling: compact polymer layer with binding capacity of approximately one protein monolayer. Suitable for various applications, including protein-protein interaction analysis.



ProteOn GLM sensor chip. For general amine coupling: extended polymer matrix with intermediate binding capacity for high response. Suitable for various applications, including protein-small molecule and protein-protein interaction analysis.



ProteOn GLH sensor chip. For general amine coupling: highly extended polymer layer for maximum binding capacity. Suitable for protein-small molecule and protein-protein interactions where highest sensitivity is the primary objective.



ProteOn NLC sensor chip. For binding of biotinylated molecules: NeutrAvidin immobilized to GLC layer. Suitable for various applications, including protein-DNA and protein-protein interaction analysis.

2. Optimize Experimental Conditions

A range of ProteOn protocol development kits and reagents are available to optimize coupling, immobilization, regeneration, and protocol development. The kits provide sufficient reagents and sensor chips to perform and analyze a complete ProteOn experiment. New users will gain familiarity with instrument setup, operation, experimental design, and data analysis. Experienced users will find the kits useful for system benchmarking and to provide positive control reagents for protocol development.



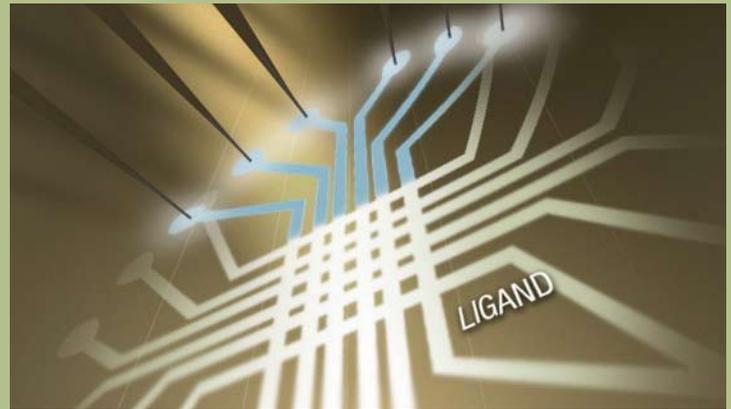
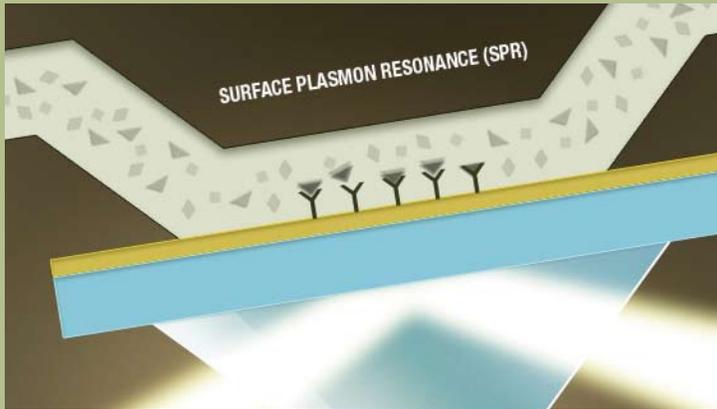
ProteOn Amine Coupling Kit



ProteOn One-shot Kinetics Kit



ProteOn Regeneration Kit



3. Create a Protocol

Create a protocol using ProteOn Manager software, a user-guided software interface for instrument control, experimental design, data collection, and analysis. The ProteOn Manager software workflow provides a guided, sequential yet flexible approach to experimental design. Get started by opening the protocol database and selecting from predefined protocol templates, or create your own custom protocol. ProteOn

Manager software lets you define your sample panels, import sample lists, or go straight to defining your protocol steps (and define your sample panels later). For convenience, ProteOn Manager software automatically associates samples to protocol steps. Additionally, ProteOn Manager software provides an easy-to-use guide for sensor chip orientation, so you always know the exact position of each sample on the 6 x 6 array.

ProteOn One Shot Kinetics Kit

Protocol Editor

Group
Click and Drag

- Immobilization
- Stabilization
- Interaction
- EVC calibration

Protocol Steps

- Initial Rack [1]
- Settings-1
- Set Temperature-25
- Buffer A
- Load EDAC/Sulfo-NHS
- Immobilization-1
 - EDAC/Sulfo-NHS
 - IL-2 Antibody
 - Ethanolamine.HCl
- Stabilization-1
- PBST,Blank
- Interaction-1
 - Add Fresh IL-2
 - PBST, Dbl Ref
 - IL-2

Step
Click and Drag

- Activate
- Ligand
- Deactivate
- Regenerate
- Analyte
- Blank
- CoInject Ligand
- CoInject Analyte
- Pause
- Set Temperature
- Set Buffer
- Change Rack

Rack-1

ProteOn One Shot Kinetic Kit Reagents

[1]: L1-L6	EDAC/Sulfo-NHS	Req. Vol. (ul)	98
Where Used	5. EDAC/Sulfo-NHS	Type	Activator
[1]: K1-K6	IL-2 Antibody	Req. Vol. (ul)	196
Where Used	6. IL-2 Antibody	Type	Ligand

Location	Sample Name	Concentration	Unit	MW (Da)	Type
K1	ProteOn IL-2 Antib...	0.00625	m...		Undefined
K2	ProteOn IL-2 Antib...	0.00625	m...		Undefined
K3	ProteOn IL-2 Antib...	0.00625	m...		Undefined
K4	ProteOn PBST, pH ...	0	m...		Blank
K5	ProteOn IL-2 Antib...	0.00625	m...		Undefined
K6	ProteOn IL-2 Antib...	0.00625	m...		Undefined

[1]: J1-J6	Ethanolamine.HCl	Req. Vol. (ul)	
Where Used	7. Ethanolamine.HCl	Type	Deactivator
[1]: I1-I6	H3PO4	Req. Vol. (ul)	244
Where Used	8. H3PO4	Type	Regenerator

[1]: C1-C6	IL-2	Concentration
1	C1	IL-2, 80 nM
2	C2	IL-2, 40 nM
3	C3	IL-2, 20 nM
4	C4	IL-2, 10 nM
5	C5	IL-2, 5 nM
6	C6	IL-2, 2.5 nM

Ligand History	Concentration
1 G1	ProteOn IL-2 Antibody/Acetate pH 4.5 fr... 0.025 mg/ml
2 G2	ProteOn IL-2 Antibody/Acetate pH 4.5 fr... 0.025 mg/ml
3 G3	ProteOn IL-2 Antibody/Acetate pH 4.5 fr... 0.025 mg/ml
4 G4	ProteOn IL-2 Antibody/Acetate pH 4.5 fr... 0.025 mg/ml
5 G5	ProteOn IL-2 Antibody/Acetate pH 4.5 fr... 0.025 mg/ml
6 G6	ProteOn IL-2 Antibody/Acetate pH 4.5 fr... 0.025 mg/ml

Step Type: **Analyte** Step Name: **IL-2** [Advanced](#)

Flow Rate: 30 μ /min \times Contact Time: 200 s = Volume: 100 μ

Dispersion: 600 s

Ordering Information

Catalog #	Description
ProteOn System, Software, and Regulatory Tools	
176-0100	ProteOn XPR36 Protein Interaction Array System , 100–240 V, includes ProteOn XPR36 instrument, 2 licensed copies of ProteOn Manager software, controller and display, communication cable, sample rack, rack needle holder, microplate needle holder, collection tank, choice of 2 sensor chips, one-shot kinetics kit, maintenance kit, 2 bottles of PBS/Tween running buffer, chip normalization solution, 200 sample vials, 25 microplates with standard wells, 50 sheets of microplate sealing film, instructions
176-0200	ProteOn Manager Software , for ProteOn XPR36 instrument control, experiment design, data collection, and analysis
176-0210	ProteOn Manager Software, Security Edition , 1-user license, includes 1 HASP key
176-4200	ProteOn XPR36 IQ/OQ Kit , includes ProteOn XPR36 IQ/OQ software, ProteOn Manager software, ProteOn operation qualification (OQ) kit
176-4225	ProteOn XPR36 Regulatory Tools Package , includes ProteOn Manager software, Security Edition, 1-user license, 1 HASP key, ProteOn XPR36 IQ/OQ kit

NeutrAvidin is a trademark of Pierce Biotechnology, Inc.

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Spain 34 91 590 5200 **Sweden** 08 555 12700 **Switzerland** 061 717 95 55 **Taiwan** 886 2 2578 7189 **United Kingdom** 020 8328 2000