

Selection guide

# Biacore™ systems





# Biacore™ systems

Biacore™ systems are designed to provide valuable information and high-quality interaction data for a broad range of molecular entities and scaffold types, including (but not limited to) of small molecules and fragments, antibodies, multidomain proteins and viruses. The systems are used in a wide range of applications across a range of scientific fields including: basic biological research, drug discovery and development, immunogenicity studies, vaccine development, and quality control. Biacore™ systems allow you to:

- Understand the relationship between molecular interactions and function or to confirm and validate your results from other techniques
- Screen and characterize for hits and optimize leads based on selectivity, affinity, and kinetics
- Screen, identify and characterize antibodies and proteins based on yes/no binding, affinity, and kinetics from the fastest on-rates to the slowest off-rates
- Quantitate protein by measuring the concentration of active protein with retained biological function

## Which system is right for you?



Applications	Biacore™ X100	Biacore™ 1 series	Biacore™ 8 series
Kinetics/affinity characterization	●	●	●
Kinetics/affinity screening	●	●	●
Biacore Single Cycle Kinetics (SCK)™	●	●	●
Parallel Kinetics™			●
2D Kinetics™			●
LMW interaction analysis	●	●	●
Fragment screening		●	●
Epitope binning	○	●	●
Immunogenicity	○	○	○
Concentration analysis	●	●	●
Calibration-free concentration analysis	●		
Parallel line analysis (PLA)		●	●
Buffer scouting using ABA command		●	●
Multi-complex analysis using Poly command		●	
GxP support		●	●
Built-in knowledge library	●		

● Application may be performed using the indicated product.

○ Application may be performed, but with limitations in software and/or hardware functionality.

## Specifications\*

	Biacore™ X100	Biacore™ 1K	Biacore™ 1K+	Biacore™ 1S+	Biacore™ 8K	Biacore™ 8K+
Association rate ( $k_a$ )	Proteins: up to $10^8 \text{ M}^{-1}\text{s}^{-1}$ LMW molecules: up to $10^6 \text{ M}^{-1}\text{s}^{-1}$	Proteins: up to $3 \times 10^9 \text{ M}^{-1}\text{s}^{-1}$ LMW molecules: up to $5 \times 10^7 \text{ M}^{-1}\text{s}^{-1}$			Proteins: up to $10^9 \text{ M}^{-1}\text{s}^{-1}$ LMW molecules: up to $10^7 \text{ M}^{-1}\text{s}^{-1}$	
Dissociation rate ( $k_d$ )	$10^{-5}$ to $0.1 \text{ s}^{-1}$	$10^{-6}$ to $1 \text{ s}^{-1}$		$10^{-5}$ to $6 \text{ s}^{-1}$	$10^{-6}$ to $0.5 \text{ s}^{-1}$	
Affinity range	pM to mM	fM to mM			fM to mM	
Concentration limit of detection (LOD)	10 pM	$\geq 1 \text{ pM}$			$\geq 1 \text{ pM}$	
Precision (concentration analysis)	< 5% CV	< 5% CV			< 5% CV	
Molecular weight limit	$M_r > 100 \text{ Da}$	No lower limit for organic molecules			No lower limit for organic molecules	
Short term noise	< 0.1 RU (RMS)	Typically < 0.03 RU (RMS)		Typically < 0.01 RU (RMS)	Typically < 0.02 RU (RMS)	
Baseline drift	< 0.3 RU/min	Typically < 0.3 RU/min			Typically < 0.3 RU/min	
Blank subtracted drift	N/A†	$< \pm 0.003 \text{ RU/min}$			$< \pm 0.03 \text{ RU/min}$	
Sample consumption	Injection volume 2 to 100 $\mu\text{L}$ plus 25 to 30 $\mu\text{L}$ (application dependent)	Injection volume 1 to 400 $\mu\text{L}$ plus 20 to 40 $\mu\text{L}$ (application dependent)			Injection volume 1 to 200 $\mu\text{L}$ plus 20 to 50 $\mu\text{L}$ (application dependent)	
Immobilized molecule consumption	Typically 1 $\mu\text{g}$	0.03 to 3 $\mu\text{g}/\text{flow cell}$			Typically 0.03 to 3 $\mu\text{g}/\text{flow cell}$	
Analysis temperature	Ambient $4^\circ\text{C}$ to $40^\circ\text{C}$ ‡	$25^\circ\text{C}$ to $37^\circ\text{C}$		$4^\circ\text{C}$ to $40^\circ\text{C}$ (at least $20^\circ\text{C}$ below ambient temperature)	$4^\circ\text{C}$ to $40^\circ\text{C}$ (maximum $20^\circ\text{C}$ below ambient temperature)	
Sample storage temperature	Ambient	$4^\circ\text{C}$ to $37^\circ\text{C}$ (at least $18^\circ\text{C}$ below ambient temperature)		$4^\circ\text{C}$ to $40^\circ\text{C}$ (at least $18^\circ\text{C}$ below ambient temperature)	$4^\circ\text{C}$ to $40^\circ\text{C}$ (maximum $18^\circ\text{C}$ below ambient temperature)	
Data collection rate	1 Hz	1 or 10 Hz		1, 10, or 40Hz	1 or 10 Hz	
Sample capacity	15 vials	1 × 96- or 384-well microplate, normal and deep-well  1 reagent rack with 21–43 positions compatible with 0.7–4.4 mL vials	2 × 96- or 384-well microplates, normal and deep-well  2 reagent racks with 21–43 positions compatible with 0.7–4.4 mL vials		4 × 96 or 384-well microplates normal and deep-well	12 × 96- or 384-well microplates normal and deep-well
Number of flow cells	2 in 1 channel	6 in 1 channel			16 in 8 channels	
Flow cell addressing	Single, pairwise	Single, pairwise	Single, pairwise, serial up to six			Single, pairwise
Unattended run time	24 h	60 h	72 h		60 h	72 h
Buffer selector	No	No	Yes			Yes
Method queue	No	Yes	Yes			Yes
Software	Biacore™ X100 Control and Evaluation Software	Biacore™ Insight Software				
Software extensions	Biacore™ X100 Plus Package	Extended Screening   Concentration and Potency   Epitope Binning   Data Integration   GxP   Biacore Intelligent Analysis™				
Dimensions (W × H × D)	596 × 563 × 593 mm	755 × 725 × 666 mm			902 × 875 × 616 mm	

\* Specifications are representative values, which can vary dependent on experimental conditions and individual properties of ligand and analyte.

† Not determined on Biacore™ systems released before 2016.

‡ Biacore™ X100 Plus Package

# Label-free interaction analysis – apply to your research, drug discovery and development, and QC



## Biacore™ X100

### Your research – boosted.

Robust two flow-cell SPR system for reliable insights into biological processes in multiuser environments.

- One needle SPR system for small-scale interaction analysis.
- Characterize molecular mechanisms and interaction pathways based on kinetics and affinity.
- Gain increased understanding of structure-function relationships.
- Determine the active concentration without the need for standard curves.
- Guided workflows for novice users with easy access to a knowledge library.

## Biacore™ 1K

### Your research – streamlined.

Six flow-cell SPR system for robust, reproducible interaction analysis from day one with minimal effort. Affordable cost and reduced running expenses.

- No programming skills needed to design, set up or start your assay.
- Fast and flexible exploratory studies using Interactive run. Take full control of the instrument and make decisions based on your results.
- The one needle SPR system with novel injection tool enables multicomplex analysis in one run.
- Upgradable to Biacore™ 1K+ for higher capacity and increased analytical flexibility.

## Biacore™ 1K+

### Your development – advanced.

Six flow-cell SPR system with full flexibility for assay design and increased capacity for interaction analysis, with straightforward transfer of methods between systems.

- Serial injection over all six flow cells reduces analyte consumption by up to 40% for a small antibody screen, compared to a 4 flow-cell SPR system.
- The one needle SPR system tackles complex questions and samples. Queue up multiple methods, use buffer selector and the innovative sample injection tools to get quality data and results.
- Simpler data interpretation: compile, visualize and export data with results in minutes.
- Optimized for use in GxP regulated environment.

## Biacore™ 1S+

### Your discoveries – elevated.

Six flow-cell SPR system with exceptional sensitivity, full flexibility, and high capacity for interaction analysis of most challenging targets or unstable binders with low response level.

- The one needle SPR system with the highest sensitivity and data collection frequency allows you to better distinguish tight binders.
- Broad analysis temperature range for expanded characterization and stability studies.
- Increase operational efficiency, line up methods, utilize the sample capacity for unattended runtime of 72 h.
- Optimized injection design reduces sample consumption by 13% compared to a typical kinetic experiment.

## Biacore™ 8K

### Your discovery – accelerated.

16 flow-cell SPR system that efficiently delivers binding data with outstanding quality and high capacity, meeting your toughest challenges in screening, characterization, process optimization, and quality control.

- High sample capacity of up to 4 × 384-well microplates.
- 60 h unattended run time with queueing abilities and rapid multi-run evaluations; option for analysis in a GxP-regulated environment.
- Eight needle SPR system for interaction analysis allowing ranking, kinetics, affinity, epitope binning, concentration, and relative potency.
- Superb data quality while increasing your throughput 8 times vs one-needle SPR systems.
- Parallel analysis enables rapid optimization of assay conditions and troubleshooting.

## Biacore™ 8K+

### Your discovery – accelerated. With maximized capacity.

16 flow-cell SPR system that efficiently delivers binding data with outstanding quality and maximized capacity, meeting your toughest challenges in screening, characterization, process optimization, and quality control.

- Exceptional sample capacity of up to 12 × 384-well microplates.
- 72 h unattended run time with queueing abilities and rapid multi-run evaluation; option for analysis in a GxP-regulated environment.
- Eight needle SPR system for interaction analysis allowing ranking, kinetics, affinity, epitope binning, concentration, and relative potency.
- Superb data quality while increasing your throughput 8 times vs one-needle SPR systems.
- Parallel analysis enables rapid optimization of assay conditions and troubleshooting.

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