

Single-Use RED Plate with Inserts

MAN0011619

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Number	Description
90006	Single-Use RED Plate with Inserts, 8K MWCO, 1 each
90007	Single-Use RED Plate with Inserts, 8K MWCO, 5 each
99006	Single-Use RED Plate with Inserts, 8K MWCO, 10 each
90112	Single-Use RED Plate with Inserts, 12K MWCO, 1 each
91012	Single-Use RED Plate with Inserts, 12K MWCO, 10 each

Storage: Upon receipt store inserts at room temperature.

Introduction

The Thermo Scientific™ Single-Use RED (rapid equilibrium dialysis) Plate is composed of disposable high-density polypropylene and is pre-loaded with 48 equilibrium dialysis membrane inserts. Each insert is comprised of two side-by-side chambers separated by an O-ring-sealed vertical cylinder of dialysis membrane with varying molecular-weight cutoffs (MWCO). This pre-loaded disposable device is automation-friendly, providing operation convenience for scientists conducting protein binding applications. No pre-conditioning of the membrane inserts is needed. When using radioactive materials, this single-use plate is easily disposed of to avoid contamination and cleaning. RED Inserts and Base Plates are also available separately (see Related Thermo Scientific Products Section).

Equilibrium dialysis is an accurate and reliable method for determining protein binding affinities to chemical or biological substances of low molecular weight (see our website for detailed information on equilibrium dialysis). The RED Device is specifically designed and extensively validated for plasma serum binding assays and produce results consistent with those reported in the literature (see Appendix). In addition to plasma protein binding, the device is used for determining drug partition between red blood cell and plasma; protein binding of liver microsomes to improve the correlation between *in vitro* and *in vivo* intrinsic clearance; and the competition between tissue protein binding against plasma proteins. The Single-Use RED Plate is validated for minimal nonspecific binding.

The design of the Single-Use RED Plate provides many advantages. This format requires no extensive assembly steps or specialized equipment, and each chamber/well is easily accessible from the top of the device. The base plate has a standard 96-well plate footprint with 9 × 9mm well spacing. Additionally, the high membrane surface-to-volume ratio allows rapid dialysis, where equilibrium can be reached in 4 hours with high levels of reproducibility and accuracy.

Additional Materials Required

- Dialysis buffer: for example, phosphate-buffered saline (PBS) containing 100mM sodium phosphate and 150mM sodium chloride (Product No. 28372)
- Sealing Tape for 96-Well Plates (Product No. 15036)

Procedure for Equilibrium Dialysis

The Single-Use RED Plate is supplied ready to use for dialysis by simply adding the plasma/compound and buffer to the corresponding chambers. There is no need to pre-condition the membrane inserts. The following is an example protocol, and specific applications and analysis methods might require optimization.

1. For each replicate, prepare samples (50-500µL) by spiking test compounds with plasma or serum at the appropriate concentrations. To minimize potential errors during sample processing, perform tests in triplicate.
2. Place 50-500µL of sample into the sample chamber, which is indicated by the colored retainer ring.
3. Add a volume of dialysis buffer to the buffer chamber relative to sample used as indicated in the table below. Using the appropriate amount of buffer is essential to avoid sample volume changes.

<u>Sample Chamber</u>	<u>Buffer Chamber</u>
50µL	300µL
100µL	350µL
200µL	400µL
300µL	550µL
400µL	600µL
500µL	750µL

4. Cover the unit with sealing tape and incubate at 37°C on an orbital shaker at approximately 250 rpm or 20 rpm on an up-and-down shaker. Generally, incubating for 4 hours is sufficient to achieve equilibrium; however, actual time required might differ depending on the test compounds and shaker used. For best results, before processing samples perform a test run to empirically determine the time required to reach equilibrium.

Alternative 100-120 minute procedure: Use an agitation device such as a vortex mixer or shaker that can secure the deep-well plate. Set the mixer at ~800 rpm or the shaker at 300 rpm.

Note: Times greater than 4 hours can be used; however, an excessively long incubation (≥ 18 hours) could promote compound instability or result in a sample volume increase from hydrostatic pressure.

5. Remove seal. Minimal to no volume change should have occurred.
6. Pipette equal volumes from both the buffer and the plasma chambers and place in separate microcentrifuge tubes or into a deep-well plate for analysis. Follow the desired sample preparation procedure for sample analysis.

Procedure for Sample Analysis

Determine the test compound concentrations in the plasma and buffer samples to determine percent bound. Alternatively, compare area ratios against an internal standard between the buffer sample and plasma sample to obtain unbound drug fractions. Some common analysis methods include LC/MS/MS, radioactivity and UV/visible/fluorescent spectrometry. The following example protocol is for analysis by LC/MS/MS and may be modified as needed.

1. Pipette 25µL (if the sample used is a 50µL volume) or 50µL each of post-dialysis samples from the buffer and the plasma chambers into separate microcentrifuge tubes or plate.
2. Add a corresponding 25µL or 50µL of plasma to the buffer sample and an equal volume of buffer to the collected plasma sample.
3. Add 300µL of Internal Standard containing precipitation buffer (such as cold 90/10 acetonitrile/water with 0.1% formic acid) to precipitate protein and release compound. Vortex and incubate 30 minutes on ice.
4. Centrifuge for 10 minutes at 13,000-15,000 × g.
5. Transfer supernatant to a vial or plate for analysis. Alternatively, dry the supernatant and reconstitute before LC/MS/MS.
6. Determine the test compound concentration in the buffer and plasma chambers from peak areas relative to the internal standard. Calculate the percentage of the test compound bound as follows:

$$\begin{aligned} \% \text{ Free} &= (\text{Concentration buffer chamber} / \text{Concentration plasma chamber}) \times 100\% \\ \% \text{ Bound} &= 100\% - \% \text{ Free} \end{aligned}$$

Appendix

A. Data Comparison

The bound drug (%) in human plasma measured using the Single-Use RED Plate on high, medium and low protein-binding compounds were similar to values reported in the literature. Compounds were tested at 1 μ M.

Compound	Human Plasma (% bound)	
	RED Plate	Other Device*
Warfarin	99	99
Taxol	96	95-98
Propranolol	92	80-92
Vinblastine	99	99
Verapamil	90	88-92
Atenolol	4	< 5
Antipyrine	0	0

*Values reported in the literature.¹⁻⁵

Related Products

90085	Competition RED Base Plate , contains base plate body and lid, 1 unit
90087	Competition RED Inserts , contains eight dual- and two single-membrane inserts, 10/pkg
90036	Protein Precipitation Plates , 2 plates
89809	RED Device Inserts , 50 inserts
89810	RED Device Inserts , 250 inserts
89811	Base Plate (made of Teflon™ Material)
90004	Single-Use RED Base Plate (empty) , 2 plates
90005	Single-Use RED Base Plate (empty) , 10 plates
28372	BupH™ Phosphate Buffered Saline Packs , 40 packs
15036	Sealing Tape for 96-Well Plates , 100/pkg

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5. Steele, W., *et al.* (1983). The protein binding of vinblastine in the serum of normal subjects and patients with Hodgkin's disease. *Eur J Clin Pharmacol* **24**:683-7.

General References

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