Version 6.01.10

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HRP-Antibody All-In-One Large Scale Conjugation Kit

Technical Manual

Catalog # A-9302-001

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- Check regularly for chemical leaks or spills. If a leak or spill occurs, follow the manufacturer's clean-up procedures as recommended in the MSDS.
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Chapter 1: Introduction

a. User Manual

This manual provides instructions for using the HRP-Antibody All-in-One Large Scale Conjugation Kit.

This chapter contains the following sections:

Purpose of Manual Intended Users Customer Service and Technical Support

b. Purpose of Manual

The purpose of this manual is to provide the user with the necessary instructions and reagents to produce one (1) HRP-antibody conjugate.

Use of the kit results in:

- The modification of a user-supplied IgG antibody (5 milligrams) using S-HyNic linker.
- The conjugation of HyNic-modified IgG with 4FB-HRP resulting in formation of an HRPantibody conjugate.
- Spin filter isolation of highly purified IgG-HRP conjugate; yielding approximately 4.5 milligrams of material free of residual enzyme and antibody.

c. Intended Users

The HRP-Antibody All-in-One Kit is designed for users with minimal or no conjugation experience allowing them to prepare customized, high purity, ready-to-use HRP-antibody conjugates within a single day.

d. Customer Service and Technical Support

Additional technical information can be found at:

Telephone	Email
1-888-625-0670 (Toll Free)	Solulink@Solulink.com
Fax	Address
1-858-625-0770	Solulink-The Conjugation Company
	9853 Pacific Heights Blvd, Ste H
	San Diego, CA 92121



Chapter 2: Overview of Conjugation

a. Product Description

Each HRP-Antibody All-in-One Conjugation Kit provides all the necessary components to produce one (1) highly purified HRP-IgG conjugate. The kit requires a user to provide 5 milligrams of starting IgG antibody. Any suitably purified monoclonal or polyclonal antibody regardless of species origin or IgG subclass can be conjugated and purified within 5 hours (60 minutes hands-on). The components of this kit feature a stable, high-activity, pre-activated horseradish peroxidase (>250U/mg) and a novel Q spin filter that delivers purified conjugate in high yield (approx. 4.5 mg). Conjugates are > 95% pure, free of both residual antibody and un-conjugated HRP. Conjugates produced are guaranteed to provide optimum signal to noise in sensitive downstream applications.

All-in-One conjugation kits are based on Solulink's patented HydraLinkTM chemistry. This chemistry relies on a specific reaction between an aromatic hydrazine (HyNic) and an aromatic aldehyde (4FB); leading to formation of a stable bis-arylhydrazone bond. HydraLinKTM conjugation chemistry is capable of efficiently converting nearly 100% of an antibody to its conjugate form. This efficiency is made possible through the recent discovery that aniline rapidly catalyzes hydrazone bond formation (1, 2, 3). Aniline's ability to increase both the rate and efficiency of conjugate formation under mild reaction conditions yields reproducible and quantitative conversion of free antibody to conjugate.

Complete conversion of conjugate greatly simplifies downstream purification. Conjugate purification consists of selectively binding conjugate to a novel Q spin filter membrane that allows excess HRP to flow through unbound. This spin filter provides high purity without sacrificing yield. Conjugates made with these kits are compatible with all downstream applications requiring high immunological specificity and sensitivity such as Westerns, ELISAs, or IHC. Finally, each kit yields between 4 and 5 mg of highly purified HRP-antibody conjugate.

b. All-in-One Technology

1) Conjugation Chemistry

HydraLinK[™] chemistry is based on the use of two complementary heterobifunctional linkers; **S-HyNic** and **Sulfo-S-4FB** (Figure 1). **S-HyNic** (Succinimidly-6-hydrazinonicotinamide) is used to modify and incorporate protected aromatic hydrazines (HyNic groups) into the antibody via acylation of lysine residues. In a similar fashion a second linker known as **Sulfo-S-4FB** (Sulfo-N-succinimidly-4-formylbenzamide) is used at Solulink to form a pre-activated, high activity form of HRP called 4FB-HRP (provided in the kit). Incubation of HyNic-modified antibody with pre-activated 4FB-HRP in the presence of aniline catalyst



leads to rapid and efficient conversion of antibody to conjugate through formation of stable bis-arylhydrazone bonds (Figure 2).



Figure 1. Structure of S-HyNic and Sulfo-S-4FB linkers used in conjugating HRP to antibody.



Figure 2. Aniline catalyzed formation of IgG-HRP conjugate.



2) Conjugate Purification

The efficiency of catalyzed hydrazone bond formation greatly simplifies conjugate purification. Because aniline increases both the rate and efficiency of conjugate formation under mild reaction conditions; it leads to near quantitative (>97%) conversion of free antibody to conjugate; leaving behind only excess 4FB-HRP. Purification then simply consists of selectively binding the conjugate to a novel filter (Q spin filter) that exploits biophysical properties of the antibody portion of the conjugate (4, 5) while permitting free HRP to flow through unbound. In this manner conjugate is eluted in highly purified form with high yield (4 to 5 milligrams).



Figure 3. Q spin filter purification of HRP-IgG conjugates.





c. All-in-One Conjugation Process Summary



d. Materials Provided and Storage Conditions

Components	Amount	Storage conditions
S-HyNic	1 x 500 μg	Keep refrigerated within desiccated
		sealed aluminum pouch
4FB-Modified HRP	2 x 1.1 ml	Keep refrigerated (2-8°C)
Buffer A	5 ml	Keep refrigerated (2-8°C)
Buffer B	5 ml	Keep refrigerated (2-8°C)
Buffer C	30 ml	Keep refrigerated (2-8°C)
Buffer D	1	Keep refrigerated (2-8°C)
PBS	5 ml	Keep refrigerated (2-8°C)
5ml Spin Column	5	Keep refrigerated (2-8°C)
Q Spin Filter	1	Room temperature or refrigerated
Q Collection Tubes	2	Room temperature or refrigerated
DMF	0.5 ml	Room temperature or refrigerated
Diafiltration Spin Filters	1	Room temperature or refrigerated

e. Additional Materials Required But Not Provided

Bradford Protein Assay Reagents (verification of final conjugate concentration) Standard UV-VIS or NanoDrop[™] Spectrophotometer Pipettes (P-10, P-100, P-1000) and tips Table Top Centrifuge (e.g. variable speed capable of handling 15 and 50 ml disposable tubes)

1.5 microfuge tubes, 15 and 50 ml disposable tubes

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Chapter 3: Antibody-HRP All-in-One Conjugation Protocol

a. IgG Sample Preparation

Antibodies come in two physical forms, solids or liquids. Individual samples can vary significantly in the amount of packaged IgG (protein mass) and/or concentration (mg/ml). We highly recommend that IgG concentrations be confirmed either by a Bradford protein assay or A280 whenever possible. The All-in-One conjugation protocol requires that antibody samples be free of protein carriers such as BSA or gelatin before proceeding. A 5 milligram quantity of antibody is required to start the procedure. Depending on the state of your initial sample (solid or liquid), proceed as follows:

Antibody is in Solid Form (e.g. lyophilized powder)

Resuspend the lyophilized antibody (5 mg free of protein additives gelatin or BSA) in 1.25 ml **Buffer A** to obtain a 4 mg/ml solution. If the antibody sample contains less than 5 mg per vial (e.g. 1 mg), resuspend the requisite number of vials equivalent to 5 mg in 1.25 ml **Buffer A** to obtain a 4 mg/ml solution.

Proceed to step b.

Antibody is in Liquid Form (e.g. PBS or TBS Buffer)

If an antibody sample is in liquid form at 4 mg/ml, simply transfer 1.25 ml to a labeled microfuge tube (5 mg). If the sample is in liquid form at a concentration greater than 4 mg/ml, transfer a volume equivalent to 5 mg antibody to a labeled microfuge tube and add **Buffer A** to obtain a 4 mg/ml solution. If a sample is at a concentration less than 4 mg/ml, concentrate the sample to 1.25 ml and 4 mg/ml using any suitable ultra-filtration spin filter (e.g. Amicon or VivaSpin) as described in the Appendix. A concentration filter is provided with this kit.

Proceed to step b.

b. Buffer Exchange IgG

- 1. Prepare a 5 ml spin column (provided) by twisting off the bottom closure and loosening the cap (do not remove). Place the spin column into a 15 ml collection tube (not provided).
- 2. Mark the top of the cap using an indelible pen to identify the sample and place a vertical mark on the side of each spin column as illustrated on the next page.





5 ml spin column and 15 ml collection tube

- 3. Place the assembly into the table top centrifuge and orient the vertical mark on the spin column aiming outward and away from the center of the rotor.
- 4. Centrifuge at **1,000 x g for 2 minute**. Discard the flow through from the collection tube. The column matrix will appear white in color. Place the column back into the empty 15 ml collection tube.
- 5. Add 2.5 ml of **Buffer A** to the top of dry resin bed. Centrifuge at **1,000 x g for 2 minutes**. Discard the flow through from the collection tube.
- 6. Repeat step 5 two (2) additional times. Place the column back into a **new empty 15 ml collection tube (not provided).**
- 7. Remove the cap; load the antibody sample (1.25 ml at 4 mg/ml) to the top of the dry resin bed; loosely recap and place the column back into the collection tube.
- 8. Orient the spin column mark outward as before and centrifuge at **1,000 x g for 2 minutes**. Use an appropriate balance tube opposite the assembly.

IMPORTANT: rotor speed should be set to $1000 \times g$ (**RCF**) and not $1000 \times rpm$ (**RPM**). The volume recovered should always be approximately the same volume that was loaded on the spin column (e.g. 1.25 ± 0.2 ml). If the recovered volume is low, the centrifuge may require recalibration. If volume is low; re-centrifuge at the appropriate speed in an attempt to recover the full volume (i.e. 1.25 ml).

9. Cap and label the 15 ml collection tube containing buffer exchanged antibody.

c. HyNic Modify IgG

1. Add 100 μ l DMF to S-HyNic reagent vial. Pipette the solution up and down to resuspend the reagent pellet.



- 2. Add 58 μ l dissolved S-HyNic reagent to the antibody solution (1.25 ml @ 4mg/ml). Pipette the solution up and down to mix.
- 3. Incubate the reaction for 2 h at room temperature. Set the mixture in a dark place or cover with aluminum foil.

d. Buffer Exchange IgG

- 1. Ten minutes before the end of the IgG modification reaction prepare a 5 ml spin column (provided) by twisting off the bottom closure and loosening the cap (do not remove). Place the spin column into a 15 ml collection tube (not provided).
- 2. Mark the top of the cap using an indelible pen to identify the sample and place a vertical mark on the side of each spin column as shown below.



5 ml spin column and 15 ml collection tube

- 3. Place the assembly into the table top centrifuge and orient the vertical mark on the spin column aiming outward and away from the center of the rotor.
- 4. Centrifuge at **1,000 x g for 2 minute**. Discard the flow through from the collection tube. The column matrix will appear white in color. Place the column back into the same empty 15 ml collection tube.
- 5. Add 2.5 ml of **Buffer B** to the top of dry resin bed. Centrifuge at **1,000 x g for 2 minute**. Discard the flow through from the collection tube.
- 6. Repeat step 5 two (2) additional times. Place the column back into a **new** empty 15 ml collection tube (not provided).



- Remove the cap; load the entire antibody/HyNic modification reaction (~1.25 ml at 4 mg/ml) to the top of the dry resin bed; loosely recap and place the column back into the collection tube.
- 8. Orient the spin column mark outward as before and centrifuge at **1,000 x g for 2 minutes**. Use an appropriate balance tube opposite the assembly.
- 9. Cap and label the 15 ml collection tube containing the buffer exchanged antibody.

e. Conjugate Formation

- 1. Spin two dark brown vials containing 4FB-modified HRP (5 seconds @ 1000 x g) to collect the contents at the bottom of each tube.
- 2. Transfer the two 1.1 ml 4FB-modified HRP aliquots to the tube containing HyNicmodified antibody (1.25 ml); pipette the mixture (~3.5 ml) up and down to mix. Set the reaction mixture in a dark place or cover with aluminum foil.
- 3. Incubate for 2 h at room temperature.

f. Buffer Exchange Conjugate

- 1. Ten minutes prior to the end of the conjugation reaction prepare two 5 ml spin columns (provided) by twisting off the bottom closure and loosening the cap (do not remove). Place the spin columns into new 15 ml collection tubes (not provided).
- 2. Mark the top of the cap using an indelible pen to identify the sample and place a vertical mark on the side of each spin column as shown below.



5 ml spin column and 15 ml collection tube

3. Place the two assemblies into a table top centrifuge and orient the vertical mark on the spin columns aiming outward and away from the center of the rotor.



- 4. Centrifuge at **1,000 x g for 2 minute**. Discard the flow through from each collection tube. The column matrix will appear white in color. Place the columns back into the same empty 15 ml collection tubes.
- 5. Add 2.5 ml of **Buffer C** to the top of each dry resin bed. Centrifuge at **1,000 x g for 2 minutes**. Discard the flow through from each the collection tube.
- 6. Repeat step 5 two (2) additional times. Place the column back into a **new** empty 15 ml collection tube (not provided).
- Remove the caps; load 1.75 ml conjugate onto each spin column (Section e, step 3) to the top of the dry resin bed; loosely recap and place the columns back into their collection tubes.
- 8. Orient the spin column mark outward as before and centrifuge at **1,000 x g for 2 minutes**.
- 9. After centrifugation, add 3.5 ml **Buffer C** to the bottom of each collection tube containing the conjugate (~5.25 ml volume in each collection tube) and pipette up and down to mix. Set both collection tubes aside containing the conjugate in a dark place or cover with aluminum foil.

g. Q Spin Filter Purification

1. Pre-wet a Q spin filter by adding 1.5 ml **Buffer C** to the top of the unit (see filter below) and incubate for 2 minutes.



- 2. Place the spin filter assembly into a table top centrifuge and orient the letter Q towards the center of the rotor; spin at **500 x g for 4 minutes**; discard the flow-through and place the filter back into the empty collection tube.
- 3. Load one-half of the antibody-HRP conjugate volume (~5.25 ml, section f step 9) to the top of the filter unit and incubate for 2 minutes.
- 4. Place the oriented and balanced assembly in the table top centrifuge and spin at **500 x g** for 4 minutes.



- 5. Load the second half of the antibody-HRP conjugate volume (~5.25 ml, section f, step 9) to the top of the same filter unit and incubate for 2 minutes.
- 6. Place the oriented assembly in the centrifuge and spin at **500 x g for 4 minutes**; discard the flow-through from the collection tube and place the Q spin filter back into the same empty collection tube. **Note-** a brown colored conjugate will be visible on the top of the Q membrane.
- Add 5 ml Buffer C to the filter unit; orient in the table top centrifuge and spin at 500 x g for 4 minutes; discard the flow-through from the bottom collection tube and place the filter back into the same empty collection tube.
- 8. Repeat step 7 three (3) additional times.
- 9. Remove the Q spin filter unit from its collection tube and place it into a new collection tube (provided).
- 10. Add 0.5 ml **Buffer D** to the top of the Q spin filter containing the tightly bound brown conjugate and incubate for 5 minutes on the bench top.
- 11. Place the oriented and balanced assembly in a table top centrifuge and spin at **500 x** g for 4 minutes to elute.
- 12. Repeat step 10 and 11 two (2) additional times. Total elution volume will be 1.5 ml. Set the collection tube aside on the bench.

h. Buffer Exchange Purified Conjugate

- 1. Prepare a 5 ml spin column (provided) by twisting off the bottom closure and loosening the cap (do not remove). Place the spin column into a 15 ml collection tube (not provided).
- 2. Mark the top of the cap using an indelible pen to identify the sample and place a vertical mark on the side of the spin column as shown below.



5 ml spin column and 15 ml collection tube



- 3. Place the balanced assembly into a table top centrifuge and orient the vertical mark on the spin column aiming outward and away from the center of the rotor.
- 4. Centrifuge at **1,000 x g for 2 minute**. Discard the flow through from the collection tube. The column matrix will appear white in color. Place the column back into the same empty 15 ml collection tube.
- 5. Add 2.5 ml of **PBS buffer** to the top of dry resin bed. Centrifuge at **1,000 x g for 2 minute**. Discard the flow through from the collection tube.
- 6. Repeat step 5 two (2) additional times.
- 7. Place the spin column into a **new** 15 ml collection tube (not provided).
- 8. Remove the cap; load 1.5 ml eluted conjugate (Section g, step 12) to the top of the dry resin bed; loosely recap and place the column back into the collection tube.
- 9. Orient the spin column mark outward as before and centrifuge at **1,000 x g for 2 minutes**. Use an appropriate balance tube opposite the assembly.
- 10. After centrifugation, transfer the purified conjugate solution (1.5 ml) from the bottom of the collection tube to a new 1.5 ml tube and label appropriately.
- 11. Measure the final conjugate protein concentration using a Bradford or BCA protein assay (see Appendix).

Chapter 4: Appendix



a. Monoclonal IgG-HRP Conjugate: An Example

Figure 5. Coomassie-stained (4-12% SDS-PAGE) gels illustrating typical conjugation results. Horseradish peroxidase is a 44 kD highly glycosylated protein that migrates as a broad, high M.W. band when the protein sample is not heated (70° C) before loading on the SDS-PAGE gel.

Lane 1.	Protein M.W. Marker
Lane 2.	4FB-HRP (1x LDS sample buffer, unheated sample 7.5 μg)
Lane 3.	4FB-HRP (1x LDS sample buffer, heat-treated 70°C 4 min 7.5 μ g)
Lane 4.	GK1.5 mAb (1x LDS sample buffer, unheated sample 1.5 μg)
Lane 5.	GK1.5 mAb (1x LDS sample buffer, heat-treated 70°C 4 min 1.5 μg)
Lane 6.	GK1.5 /HRP crude conjugation rxn (1x LDS sample buffer, unheated sample 7.5 μ g)
Lane 7.	GK1.5 /HRP purified conjugate (1x LDS sample buffer, unheated sample)
Lane 8.	GK1.5 /HRP purified conjugate (1x LDS sample buffer, heat-treated 70°C 4 min)
Lane 9.	GK1.5 /HRP Q filter 1 st flow-through 20 μ L (1x LDS sample buffer, unheated sample)
Lane 10	. GK1.5 /HRP Q filter 1 st flow-through 20 μL (1x LDS sample buffer, heat-treated 70°C 4
	min)

b. Direct ELISA Assay Using IgG-HRP All-in-One Conjugate



Figure 6. Direct *ELISA* curves generated using an HRP conjugate made with the All-in-One kit. A mouse anti-FITC monoclonal antibody was conjugated to HRP as described in the manual. Antigen consisting of FITC-labeled BSA (FITC MSR = 2) was coated on plates in a 2-fold dilution series (100 μ l @ 500, 250, 125, 62.5, 31.25, 15.625, 7.8, 3.90, and 1.95 ng/ml) using standard methods. Immobilized antigen was then detected at 3 different conjugate concentrations (1 μ g/ml. 0.5 μ g/ml. 0.25 μ g/ml) using TMB substrate (20 minutes @ 450 nm) on a Molecular Devices plate reader.



c. Bradford Protein Assay

Solulink highly recommends that whenever IgG is not limiting or its concentration, source, or quality are unknown that the sample be assayed for initial protein concentration using a Bradford protein assay prior to conjugation. The starting quality and quantity of antibody is critical to the success of the procedure. A reference assay protocol is provided for measuring antibody or conjugate protein concentrations using Bradford protein reagents (not provided in this kit).

Bradford Microtiter Plate Procedure

Required Materials

Bradford Reagent (Bio-Rad, Hercules, CA, Cat. #500-0006) 96-well microtiter plate (standard flat bottom) PBS (phosphate buffered saline) P-200 and P-1000 pipettes Bovine IgG Antibody Standard: 2 mg/ml (Pierce/ThermoFisher, Cat. # #23212) Molecular grade water

Assay Protocol

- 1) Prepare 2 ml of a Bradford working solution by adding 400 μ l dye reagent to 1600 μ l molecular grade water (1:4 ratio).
- 2) Prepare the following protein dilution standards and blank as follows:

Add 160 μ l 2 mg/ml bovine IgG standard to 240 μ l PBS (0.8 mg/ml standard) Add 150 μ l 0.8 mg/ml standard to 50 μ l PBS (0.6 mg/ml standard) Add 75 μ l 0.6 mg/ml standard to 25 μ l PBS (0.4 mg/ml standard) Add 50 μ l 0.4 mg/ml standard to 50 μ l PBS (0.2 mg/ml standard) Add 50 μ l 0.2 mg/ml standard to 50 μ l PBS (0.1 mg/ml standard) Add 50 μ l PBS (buffer blank)

- 3) Pipette 5 μ l of each standard (and blank) along with duplicates of appropriately diluted antibody sample into separate microtiter wells.
- 4) Add 100 μ l of previously diluted dye reagent (1:4) to each well and mix thoroughly. Always replace pipette tips between additions.
- 5) Incubate at room temperature for 5-10 minutes (but no more than 60 minutes).
- 6) Measure absorbance at 595 nm on a suitable microtiter plate reader.



7) A typical Bradford plate assay result from a commercial plate reader is illustrated in Figure 7.



Figure 7. Print out from a Bradford plate-based protein assay.



d. Using a NanoDrop[™] to Measure Antibody Concentration

If an antibody sample is free of protein-based carriers (e.g. BSA, gelatin) or certain interfering preservatives such as thimerosal, then a simple non-destructive scan of the IgG sample on a NanoDropTM spectrophotometer can be used to estimate antibody concentration saving the trouble of conducting a Bradford protein assay to confirm protein concentration. To estimate antibody concentration using a NanoDropTM spectrophotometer, proceed as follows.

- 1. Turn on the NanoDrop[™] spectrophotometer and click on the NanoDrop[™] icon to launch the software.
- 2. Place a 2 μ l drop of molecular grade water on the clean pedestal, click OK.
- 3. When the main menu appears, select the A280 menu option. Note- do not use the UV-VIS menu option on the NanoDrop[™] to read an antibody sample.
- 4. After the A280 menu appears, **click-off the 340 nm normalization option** using the mouse. **Note-**some instruments do not use this normalization feature in which case this step can be ignored.
- In the window labeled Sample Type, select 'Other protein E1%' option from the pulldown menu. Enter the appropriate E1% value (Table 1 on the next page) corresponding to your particular antibody sample type. For example, 14.00 for mouse IgG.
- 6. Blank the NanoDropTM spectrophotometer by placing a 2 μ l drop of the appropriate sample buffer (e.g. PBS) and click on the 'Blank' icon.
- 7. Immediately re-click the 'Measure' icon to validate the baseline (i.e. flat across the bandwidth). Clean the pedestal and repeat (if necessary) until a flat baseline is obtained. Note-sometimes air bubbles can become trapped on the pedestal during sample loading and cause baseline offsets. If necessary, remove air bubbles and rescan to insure a proper baseline.
- 8. Place a 2 μ l volume of antibody solution on the clean pedestal and click the 'Measure' icon. Wait until the spectrum (220-350 nm) appears in the window. Note-for precious or limited samples the majority of the 2 μ l aliquot can be recovered from the pedestal.
- 9. Record antibody concentration directly from the NanoDrop[™] display window [mg/ml]. Alternately, calculate the antibody concentration (manually) as illustrated on the following page.



Example: A mouse IgG sample at 1 mg/ml in PBS (100 μ l) was scanned as described and its concentration confirmed using equation #1 below.





Sample Calculation

Equation #1: [A280 /E1% value] x 10 mg/ml = protein concentration (mg/ml)

Example: Mouse IgG @ 1 mg/ml (Fig. 8) A280 reading (from scan in Figure 8) = 1.34 Antibody E1% value (Table 1) = 14.00

[A280 / E1% bovine IgG] x 10 mg/ml = protein concentration (mg/ml)

[1.34 / 14.00] x 10 mg/ml = 0.96 mg/ml

Antibody Source	Antibody E1% (1-cm path)
Human IgG	13.60
Human IgE	15.30
Rabbit IgG	13.50
Donkey IgG	15.00
Horse IgG	15.00
Mouse IgG	14.00
Rat IgG	14.00
Bovine IgG	12.40
Goat IgG	13.60

Table 1. Mass extinction coefficients (E1%) used for calculating antibody concentration. The E1% is the A280 of a 10 mg/ml solution (1-cm path length).



e. HRP Absorption Spectrum (Unmodified Horseradish peroxidase)



Figure 9. NanoDrop[™] absorption spectrum of unmodified horseradish peroxidase (220-550 nm) @ 0.66 mg/ml (sodium phosphate buffer, pH 6.0, 1 mm path length)

f. 4FB-modified HRP Absorption Spectrum



Figure 10. NanoDrop[™] absorption spectrum of 4FB-modified horseradish peroxidase (220-550 nm) @ 0.66 mg/ml (sodium phosphate buffer, pH 6.0, 1 mm path length)



g. Bovine IgG-HRP Conjugate Absorption Spectrum (All-in-One Purified)



Figure 11. NanoDrop[™] absorption spectrum of All-in-One IgG-HRP conjugate (220-550 nm) @ 0.96 mg/ml (sodium phosphate buffer, pH 6.0, 1 mm path length).

h. Concentration of Dilute Antibody Solutions

The HRP-Antibody All-in-One Conjugation protocol requires that initial antibody protein concentration be at 4 mg/ml in 1.25 ml. Many antibody vendors package at significantly more dilute concentrations (e.g. 0.25 to 2 mg/ml). In these instances, IgG samples will need to be concentrated to 4 mg/ml and 1.25 ml before proceeding. The All-in-One kit provides a diafiltration filter (M.W.C.O. 30k) for this purpose (Figure 12). Carefully follow the instructions below to avoid antibody loss or aggregation when using the filter to concentrate antibody.

Note-dilute antibody solutions require 5 milligrams of starting antibody (e.g. 2 ml @ 2.5 mg/ml).



Figure 12. Diafiltration spin filter used for concentrating dilute antibody samples prior to the start of All-in-One conjugation protocol.



Antibody Concentration Protocol

Note- the diafiltration spin filter provided is made to contain and process a maximum volume of 500 μ l. To process sufficient volume of dilute antibody solution equivalent to 5 milligrams multiple loadings and spins will be required.

- 1) Open the lid of the diafiltration spin filter device provided.
- 2) Transfer 500 μ l of dilute protein solution to the center of the filter cup.
- 3) Close the lid and orient the spin filter in the centrifuge so that the volume marker faces toward the center of the centrifuge rotor. Use an appropriate balance tube opposite the spin filter.
- 4) Centrifuge for 2 minutes @ 5,000 x g. Do not centrifuge for a longer periods of time to avoid antibody aggregation.
- 5) Open the filter unit and visually note the volume remaining. Bring the volume back in the spin filter concentrator body to 500 μ l by adding additional dilute antibody solution. Pipette the solution (500 μ l) up and down at least 20 times to fully resuspend the concentrated antibody away from the filter's surface.
- 6) Repeat steps 4 and 5 until a volume equivalent to 5 milligrams is processed.
- 7) Transfer the concentrated IgG solution (equivalent to 5 milligrams) to a new 1.5 ml microfuge tube and bring the volume to 1.25 ml using Buffer A to achieve a 4 mg/ml IgG solution. You may now proceed with the conjugation protocol.



i. Troubleshooting Guide

Problem	Possible Cause Recomn	nended Action
Poor conjugate yield	-initial antibody concentration and volume were incorrect or unknown.	-whenever possible verify the original starting antibody concentration using a Bradford protein assay or NanoDrop [™] to assure efficient conjugation. -concentrate or dilute the antibody sample to be conjugated into the required range (4-5 mg/ml and 25 μl)
Poor conjugate yield	Starting antibody concentration and volume are incorrect or unknown.	-preservatives can interfere with the accuracy of a Bradford protein assay. Remove all interfering preservatives such as thimerosal or proclin before performing a Bradford protein assay.
Poor HyNic modification	-presence of protein carrier (e.g. BSA or gelatin) is contaminating the antibody sample.	-remove and purify away all protein carriers such as BSA or gelatin using affinity chromatography or other methods
Poor HyNic modification	-improper mixing of HyNic reaction components	-make sure to properly mix the antibody- HyNic reaction mixture -use a calibrated P-10 pipette to insure accuracy of small volumes
	-presences of amine contaminants	-remove all non-protein amine contaminants such as glycine or Tris before modification
	-improper storage of S-HyNic reagent can lead to hydrolysis of this NHS	-keep and store S-HyNic sealed in the aluminum



	ester	pouch provided that contains dessicant.
	-initial antibody concentration was too low or too high.	measure the initial antibody concentration before proceeding (Bradford or NanoDrop)
		-concentrate or dilute the antibody sample into the recommend range (4-5 mg/ml and 25 µl) before proceeding
Low conjugate and/or antibody recovery	-low spin column recovery volume	-use a properly calibrated variable-speed centrifuge Follow recommended spin speed/time. Altered spin speeds can adversely compromise protein and/or volume recovery

j. Component Stability on Storage

Component	Stability	Storage Condition
Unopened Kit	1 yr	Refrigerated (2-8°C)
S-HyNic	1 yr	Keep in sealed aluminum pouch provided (2-8°C).
	24 h after re-suspending S-HyNic in DMF	Room temperature
HRP-Antibody Conjugate	9 month	Refrigerated (2-8°C) in final conjugate solution.
	1 yr	Refrigerated (2-8°C) (50% glycerol)
All other kit components	1 yr	Refrigerated (2-8°C)



Catalog # A-9302-001

k. References

- **1.** *Dirksen, A., Hackeng, T., Dawson, P.,* (2007). *Nucleophilic Catalysis of Oxime and Hydrazone Reactions by Aniline. ACS Poster*
- **2.** *Dirksen, A., Hackeng, T., Dawson, P., (2006). Nucleophilic Catalysis of Oxime Ligations. Angew. Chem. Int. Ed.* 45, 7581-7584
- **3.** *Dirksen, A., Dirksen, S., Hackeng, T., Dawson, P., (2006). Nucleophilic Catalysis of Hydrazone Formation and Transimination: Implications for Dynamic Covalent Chemistry. JIAICIS Communications.*
- **4.** *Lim,S., Manusu, H.P., Gooley, A.A., Williams, K. L., Rylatt D.B.,(1998). Purification of monoclonal antibodies from ascitic fluid using preparative electrophoresis. Journal of Chromatography A. Vol. 827, Issue 2, 11, Pages 329-335.*
- **5.** *Chiodi,F., Sidén, A., Ösby, E., (2005). Isoelectric focusing of monoclonal immunoglobulin G, A and M followed by detection with the avidin-biotin system. Electrophoresis, Vol. 6 Issue 3, 124-128.*