

Video Article

Affinity Labeling Detection of Endogenous Receptors from Zebrafish Embryos

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Abstract

By combining the powers of **Affinity Labeling** and **Immunoprecipitation** (AFLIP), a technique for the detection of low abundance receptors in zebrafish embryos has been implemented. This technique takes advantage of the selectivity and sensitivity conferred by affinity labeling of a given receptor by its ligand with the specificity of the immunoprecipitation. We have used AFLIP to detect the type III TGF- β receptor (TGFB3), also known as betaglycan, during early zebrafish development. AFLIP was instrumental in validating the efficacy of a TGFB3 morphant zebrafish phenotype. In the first step, embryo protein extracts are prepared and used to generate ¹²⁵I-TGF- β 2-TGFB3 complexes that are purified by immunoprecipitation. Later, these complexes are covalently cross-linked and revealed using SDS-PAGE separation and autoradiography detection. This technique requires the availability of a labeled ligand for, and a specific antibody against, the receptor to be detected, and shall be easily adapted to identify any growth factor or cytokine receptor that meets these requirements.

Video Link

The video component of this article can be found at <http://www.jove.com/video/54405/>

Introduction

Specific detection of proteins expressed during embryonic development is required to validate expression profiles obtained by measuring their cognate mRNAs with RT-PCR or *in situ* hybridization (ISH). This is commonly achieved by a western blot of embryo extracts followed by detection with specific antibodies. However, this approach is hard to apply to proteins that are in very low abundance, or that have properties that hamper their quantitative transfer during their blotting. Betaglycan, also known as the type III transforming growth factor β (TGF- β) receptor (TGFB3), is an example of these difficulties. TGFB3 is a part time membrane proteoglycan that binds TGF- β through its core protein¹, with notably higher affinity for the isoform TGF- β 2, a property that distinguishes it from any other TGF- β binding protein². TGFB3 in the zebrafish is expressed from 8 hpf on, reaching a maximum by 72 hpf, as detected by RT-PCR of its mRNA³.

However, despite the availability of a very specific antibody³, every attempt to detect its translated product by western blot proved unsuccessful. Reckoning that TGFB3's proteoglycan nature, as well as putative low abundance may be accountable for this failure, a detection method, AFLIP, which takes advantage of TGFB3 high affinity for TGF- β 2 was devised. In this method a protein extract from pooled embryos is allowed to specifically bind ¹²⁵I-labeled TGF- β 2 and the receptor-ligand complexes are purified by immunoprecipitation and cross-linked before separation by SDS-PAGE. The migration patterns observed by autoradiography of the gels revealed the presence and nature of the labeled receptor species. This approach combines the ligand specificity of affinity labeling with immunoprecipitation by specific antibodies, increasing detection range, avoiding the inefficient transfer blotting of TGFB3. Due to its inherent properties, the AFLIP assay is not a quantitative assay but can be used to confidently gauge relative experimental differences in the analyzed receptor.

Protocol

All experiments carried out in animals were approved by the Committee for Laboratory Animal Care and Use of the Autonomous National University of México (UNAM), under the CICUAL-Protocol number: FLC40-14. (CICUAL: "Comité Institucional para el Cuidado y Uso de los Animales de Laboratorio del Instituto de Fisiología Celular, Universidad Nacional Autónoma de México").

1. Preparation of Embryo Protein Extract

1. Collect 100 - 200 embryos for each condition to compare (morphants vs. wild type) at desired stage, for example 72 hr post fertilization (hpf).
2. Place embryos in Petri dishes containing fish water (see **Table 1**) and manually dechorionate embryos using fine-tipped forceps. Avoid using pronase during this step (or any other protease throughout the protocol) as it may digest the targeted receptor.
3. Place embryos in a 1.5 ml microfuge tube and wash embryos twice with 1x phosphate buffer solution (PBS) (see **Table 1**).
4. Add 500 μ l of deydolk buffer (see **Table 1**).

5. Release most of yolk by gently pipetting up and down (about 30 - 40 times) the embryos in the solution. For 72 hpf and 48 hpf embryos use blue and yellow pipette tips, respectively. Yellow tips may need to be slightly cut to avoid destroying embryos. The successful yolk release can be checked by observing embryos under microscope.
6. Collect embryos by centrifugation at 600 x g for 15 sec.
7. Discard supernatant carefully by using a pipet.
8. Wash embryos twice with 500 μ l washing buffer (see **Table 1**) by gently vortexing at lowest speed.
9. Collect embryos by centrifugation at 600 x g for 15 sec.
10. Starting from here, continue the procedure at 4 °C.
11. Discard supernatant carefully by using a pipet.
12. Resuspend embryos in 350 μ l of lysis buffer (see **Table 1**) and homogenize using a plastic pestle.
13. Incubate lysed embryos 30 min with agitation on a test tube rocker at 4 °C.
14. Centrifuge lysed embryos at 11,000 x g for 15 min in order to remove insoluble debris.
15. Transfer cleared supernatant using a pipet to a fresh tube.
16. Determine total protein by Bradford protein assay⁴, or other suitable procedure. If Bradford assay is used, perform the calibration curve in the presence of equivalent concentrations of the detergents presents in the lysis buffer, as they cause an underestimation of the protein standard⁴.

2. Detection of Endogenous Receptor Protein

1. **Affinity Labeling and Immunoprecipitation (all these steps at 4 °C)**
 1. Place 400 - 500 μ g of total embryo protein in a 1.5 ml microfuge tube and dilute to 1 μ g/ μ l with buffer 1 (see **Table 1**).
Note: In order to obtain 500 μ g of total embryo protein, about 100 - 200 embryos must be initially processed. 72 hpf embryos routinely yield ~ 5 μ g of total embryo protein, which is sufficient for betaglycan detection.
 2. Add enough volume of labeled TGF- β 2 stock to reach a final concentration of 150 pM and incubate with agitation on a test tube rocker 2 hr at 4 °C. TGF- β 2 must be labeled before AFLIP is started, with ¹²⁵I by the chloramine T method as described by Cheifetz *et al.*⁵.
CAUTION: Use shielding to minimize exposure while handling ¹²⁵I labeled ligand.
 3. Add enough volume of undiluted antibody against the receptor of interest in order to reach a 1:100 dilution and continue incubation for another 2 hr at 4 °C (incubation may be O/N). This is the optimal dilution of antiserum # 31, used in this study and described elsewhere³, but depending on the quality of the antibody, a larger or smaller dilution may be the optimal.
 4. Add 50 μ l of suitable immunoglobulin binding beads (e.g., G-protein-Sepharose, which was previously equilibrated in TNTE and resuspended 1:5 of its original volume in TNTE) and incubate for 50 min with agitation on a test tube rocker at 4 °C.
 5. Recover the beads by microcentrifugation at 11,000 x g for 20 sec.
 6. Discard supernatant in an appropriate radioactive trash container.
 7. Wash the IP-beads three times with 1 ml of IP wash buffer (see **Table 1**), by vortexing and microcentrifugation at 11,000 x g for 20 sec each time.
 8. Resuspend IP-beads in 250 μ l of IP wash buffer.
 9. Add 1.5 μ l of disuccinimidyl suberate (DSS, dissolved in DMSO at 10 mg/ml). Prepare the DSS solution just before its use in this step.
 10. Incubate for 15 min at 4 °C with agitation.
 11. In order to quench the crosslinking reaction, add 500 μ l of IP wash buffer, supplemented with enough Tris-Cl pH 7.4 stock to reach 25 mM Tris-Cl. The free amino groups in Tris capture unreacted DSS.
 12. Centrifuge at 11,000 x g for 20 sec to collect IP-beads and discard supernatant.
 13. Resuspend IP-beads in 30 μ l of reducing Laemmli buffer.
 14. Boil samples 5 min at 94 °C.
 15. Optionally, analyze samples in a gamma counter using manufacturer's protocol.
2. **Sample Analysis**
 1. Subject samples to denaturing SDS-PAGE. Use polyacrylamide at the appropriate percentage for the mass of the receptor and run gel under standard procedure.
 2. Immerse gel in fixative solution (see **Table 1**) for 30 min at RT.
 3. Wash gel with distilled water for 15 min.
 4. Place gel in previously hydrated filter paper and cover with saran wrap film.
 5. Dry gel at 80 °C for 1 hr.
 6. Expose gel on a white phosphorimager screen at RT O/N.
 7. Scan exposed screen using a phosphorimager using manufacturer's protocol.

Representative Results

Figure 1 shows a representative result obtained with AFLIP. Signals in Lane 1 come from the ^{125}I -ligand covalently linked to either the zebrafish betaglycan core protein (BG core, below the 150 kDa marker) or the BG core that has been processed to its proteoglycan form by attachment of glycosaminoglycans (GAG, smear ranging from 170 kDa to the top of the gel). This pattern of migration, a sharp core protein plus a smeared proteoglycan (due to heterogeneity in the length of the GAG chains), is characteristic of the TGFBR3². Since the DSS does not covalently link every single ligand-receptor complex formed, there is free ^{125}I -ligand appearing at the migration front of the gel (^{125}I -TGF- β 2). This free ligand was bound to the receptor, and remained bound during the immunoprecipitation, but since it was not covalently attached, detached during the SDS-PAGE procedure. Nonetheless, the strength of its signal still correlates to the one given by the labeled receptor. This can be appreciated by comparing lanes 2 and 3. At 48 hpf the zebrafish embryos exhibit lower amounts of BG than 72 hpf embryos³, which can be seen by the faint signals at the GAG and BG core, that correspond to the intensity of the signals given by the free ligand appearing at the front of the gel (Lane 2). Given that antibodies against rat BG⁶ do not cross-react with the ZF BG, the antibodies when used in the AFLIP gave no signal, therefore, serving as a negative control (Lane 3). Similar negative results were obtained with the pre-immune serum of the anti-ZF BG antibody³.

Figure 2 shows how AFLIP can be used to gauge the levels of expression of a given receptor in embryos subjected to an experimental manipulation, in this case, the decrease of betaglycan due to morpholino injection³. Lane 1 shows the level of TGFBR3 in a 72 hpf wild type embryo, while Lane 2 shows the effect of administering 7 ng of a morpholino directed to the exon2-intron2 boundary of the ZF BG primary transcript. Note that the BG down regulation obtained with this morpholino is not reproduced by its mis-matched control (Lane 3). These results confirmed that the phenotype obtained with this morpholino was specific for the knockdown of the TGFBR3 as described before³.

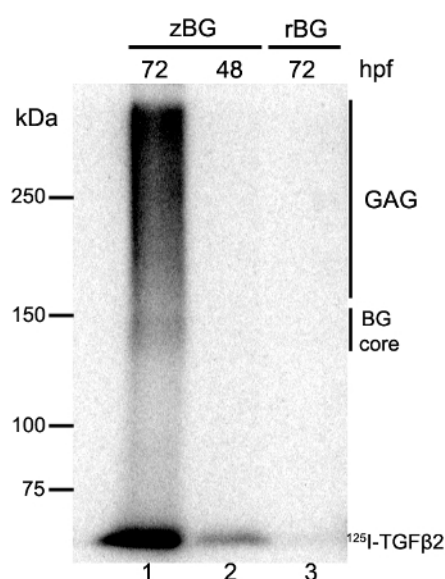


Figure 1. AFLIP Detection of BG in Zebrafish Embryos. Total protein extract from embryos at 72 hr post-fertilization (hpf) (Lanes 1 and 3) and 48 hpf (Lane 2) were subjected to ^{125}I -TGF- β 2 affinity labeling followed by IP with rabbit anti-zebrafish BG (Lanes 1 and 2, zBG) or rabbit anti-rat BG (Lane 3, rBG) as a control. In the autoradiography BG can be detected without GAG (BG core) or as a glycosaminoglycan-modified BG (GAG). [Please click here to view a larger version of this figure.](#)

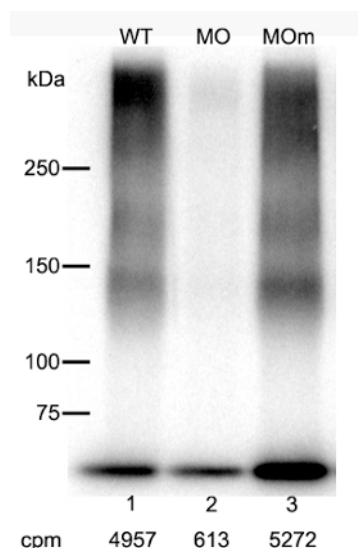


Figure 2. BG Morpholino Knockdown Detected by AFLIP. ^{125}I -TGF- β 2 affinity labeling of BG from untreated WT embryos (lane1), microinjected with specific BG morpholino (Lane 2) or a mismatch control (Lane 3). The counts per million (cpm) recovered after the immunoprecipitation and the cross-linking step are indicated. [Please click here to view a larger version of this figure.](#)

Buffer	Composition
Buffer 1	50 mM Tris-HCl pH 7.5, 150 mM NaCl, 0.1% Triton X-100
Deyolk buffer	55 mM NaCl, 1.8 mM KCl, 1.25 mM NaHCO ₃
Fish water	See Materials Table
Fixative solution	50% CH ₃ OH, 12% CH ₃ COOH, 0.185% HCHO
IP wash buffer	10 mM Na ₂ HPO ₄ , 2 mM KH ₂ PO ₄ , 137 mM NaCl, 2.7 mM KCl, 0.1% Triton X-100, 0.02% SDS, pH 7.4
Laemmli buffer	2% SDS, 10% Glycerol, 100 mM DTT, 60 mM Tris (pH 6.8), 0.001% bromphenol blue.
Lysis buffer	50 mM Tris-HCl pH7.4, 150 mM NaCl, 1 mM EDTA, 1 mM PMSF, 0.5% Triton X-100, 0.1% SDS, 0.5% Sodium Deoxycholate
PBS	137 mM NaCl, 2.7 mM KCl, 10 mM Na ₂ HPO ₄ , 2 mM KH ₂ PO ₄ , pH7.4
TNTE	50 mM Tris-HCl pH7.4, 150 mM NaCl, 1 mM EDTA, 0.1% Triton X-100
Washing buffer	110 mM NaCl, 3.5 mM KCl, 2.7 mM CaCl ₂ , 10 mM Tris-HCl pH 8.5

Table 1: Buffer Compositions

Discussion

The use of Western blots with a specific antibody against a protein of interest is a valuable tool to study its expression⁷ during embryogenesis. However, immunoblotting of highly-glycosylated proteins has not been very successful due to their inefficient transfer and weak binding to nitrocellulose or PVDF membranes^{8,9}.

Proteoglycans are a good example of this shortcoming, because of their covalently attached glycosaminoglycan chains (GAG) that are negatively charged and do not bind well to either polystyrene surfaces or hydrophobic blotting membranes. Also, the size heterogeneity of the GAG chains "spreads" the protein over a sector of the gel, effectively decreasing its amount per area of the blot. In addition, if the protein of interest has low levels of expression, its detection by regular western blots is a difficult task. The type III transforming growth factor β receptor (TGFB β R3) or betaglycan (BG), a membrane receptor with two GAG's attachment sites in mammals¹ and one in zebrafish³, has all of these properties. Although, protocols to overcome these hurdles have been devised, like the detection by avidin-biotin complex (ABC) system^{10,11} or the immunoblotting of proteoglycans on positively charged membranes¹², they cannot resolve both problems in the same protocol. Taking advantage of the high affinity of BG for its natural ligand TGF β 2 and of the availability of a specific antibody, AFLIP, a technique that combines the advantages of affinity labeling and immunoprecipitation, the above discussed limitations of western blot detection may be overcome.

Affinity labeling of membrane-bound receptors with their radiolabeled ligands is a well used tool that has been instrumental for the identification and characterization of several important growth factors receptors in cultured cells¹³⁻¹⁶. While in conventional affinity labeling a monolayer of cells is subjected to covalent ligand labeling on the tissue culture dish and then lysed, in AFLIP the ligand-receptor complexes are first formed in embryo lysates, then purified by immunoprecipitation and finally, covalently cross-linked. Because of its use of radiolabeled ligands, AFLIP is

very sensitive and in principle can be adapted to other growth factors or cytokines receptors for which a labeled ligand and a good antibody are available. This potential of AFLIP would help the detection of these low abundant but physiological relevant molecules.

Because of the inherent variability in the cross-linking step of affinity labeling, AFLIP cannot be used to quantitatively determine the levels of the measured receptors. However, if performed with adequate parallel controls, it can provide a quite accurate gauging of relative levels of their expression. As AFLIP use total protein extracts, it is not restricted to one specific cellular location of the receptor, it reveals its expression in the whole individual or organ, if applied to dissected adult specimens. Finally, if needed AFLIP could be linked to other biochemical protocols, like carbohydrate enzymatic digestions, to further characterize the receptor of interest.

Similar to conventional affinity labeling, a freshly ^{125}I -labeled ligand is strongly recommended. Given the short half-life of ^{125}I , the ligand must be used within the first 2 weeks after labeling with fresh isotope. Although most of the caveats of the protocol have been mentioned in the corresponding steps, one special mention shall be given to the careful removal of the yolk, which must be as complete and uniform as possible. Leaving uneven amounts of yolk would result in incorrect quantitation of bona fide embryo protein in the lysates.

Disclosures

The authors have nothing to disclose.

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