

Sample preparation of brain tissue for extraction of β -amyloid and other neuronal markers

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Introduction

Alzheimer's disease (AD) is characterized by regional neuronal degeneration, synaptic loss, the presence of intracellular neurofibrillary tangles (NFTs), and the progressive deposition of senile plaques in the extracellular space. While neurofibrillary tangles are composed of hyperphosphorylated Tau, the senile plaques are formed by aggregation and deposition of β -amyloid (A β) peptides. Many investigations conducted in last decade have pointed to the soluble circulating A β oligomers as the building blocks for these insoluble plaques and tangles. The increased concentration of soluble A β oligomers has also been linked to synaptic dysfunction and cognitive decline. A β peptides are a group of varying length oligomers originating through degradation of the amyloid precursor protein (APP); the two most common oligomers, A β 1-40 and A β 1-42, both have disease implications. A β 1-40 is the predominant form found in biological fluids such as blood and cerebrospinal fluid (CSF). A β 1-42 is highly hydrophobic and tends to aggregate at much lower concentrations than A β 1-40. It is the main component of plaques and thus plays a significant role in the pathogenesis of AD.

The isolation of amyloid peptides continues to present a challenge. While there are many protocols reported in the literature for the purification of amyloid peptides, a simplified, universal method remains elusive. In general, the majority of reported protocols propose a sequential approach. In first step, aqueous buffers are used to extract hydrophilic components. In the following step(s), strong detergents, acids, or strong chaotropes are applied to extract difficult to solubilize components. These steps are severely hampered by not only precipitation and protein loss but also incompatibility with many common downstream assays such as ELISA, Immunoprecipitation (IP), Western blotting (WB), or Mass Spectrometry (MS).

Here we present a study comparing relative extraction efficiency for a variety of buffers. The presented results demonstrate that, for most applications, the use of Tris-buffered Saline (TBS) for extraction of soluble A β fractions and commercial lysis buffers for extraction of "insoluble" A β fractions can effectively obviate the need for formic acid-based extraction.

Table 1. The most common protocols and buffers used for extraction of brain proteins.

Reference	Method	Buffers used
B. Kaplan <i>et. al.</i> J Clin Pathol 2003 (Review)	Sequential extractions	From TBS through guanidine hydrochloride or formic acid (FA) to MeCN/TFA
G.M. Shankar <i>et. al.</i> Nature Medicine (2008)	Sequential extractions (Ultracentrifugation)	a. TBS (20 mM Tris, 150 mM NaCl pH 7.4) b. TBS + 1% TX-100 c. TBS + 5 M guanidine chloride pH 8
A. Rostagno & J. Ghiso Curr. Protoc. Cell Biol. (2009)	Sequential extractions (Ultracentrifugation (112,000 xg 1 hr, 10°C) Centrifugation 14,000 xg Ultracentrifugation/sucrose gradient with SDS 220,000 xg/20 hr)	a. TBS b. 20 mM Tris + 2% SDS c. 70% FA d. 95% FA
E. Portelius <i>et. al.</i> Acta Neuropathol. (2010)	Sequential extractions (Centrifugation; Sonication)	a. TBS b. 70% FA
H. Schieb <i>et. al.</i> JBC (2011)	Sequential extractions (Centrifugation (16,000 xg))	a. 50 mM HEPES, 150 mM NaCl, 1% Nonidet, 0.5% Na Deoxycholate, 0.1% SDS b. PBS c. 70% FA
G.M. Shankar <i>et. al.</i> Methods Mol. Biol. (2011)	Sequential extractions (Centrifugation; Sonication)	a. TBS (15 mM Tris, 40 mM NaCl, 3 mM KCl) b. TBS with 1% TX-100 c. 88% FA (sonication) 10 mM Tris, 150 mM NaCl, 1 mM EDTA and PBS with: a. 0-60% organic solvent b. 1% detergent (also other additives) c. 1% FA
G. Shevchenko <i>et. al.</i> J. Proteome Res. (2012)	Sequential extractions (Centrifugation; Delipidation)	a. TBS (20 mM tris, 150 mM NaCl pH 7.4) b. TBS + 1% TX-100 c. 70% FA
M. Izco <i>et. al.</i> J Alzheimers Dis. (2013)	Sequential extractions (Ultracentrifugation)	a. TBS (20 mM tris, 150 mM NaCl pH 7.4) b. TBS + 1% TX-100 c. 70% FA
S. Musunuri <i>et. al.</i> J. Proteome Res. (2014)	Direct extraction (Centrifugation @ 10,000 xg; Delipidation)	10 mM Tris pH 7.4, 150 mM NaCl, 1 mM EDTA and PBS with 1% N-octyl- β -D-glucopyranoside

Materials:
Human brain tissue (Analytical Biological Services Inc.)
Grey matter from 3 non-Alzheimer's donors (healthy "H") and 3 from Alzheimer's patients ("A")

Reagents & Buffers
RIPA lysis buffer, 10X (EMD Millipore cat # 20-188)
CytoBusterTM Protein Extraction Reagent, "Cyto" (Novagen cat # 71009-50mL)
Triton X-100 Reduced, "TX-100" (Sigma-Aldrich cat # X100RS)
Triton X-114, "TX-114" (Sigma-Aldrich cat # X114)
Acetone, "MeCN" (Sigma-Aldrich cat # AX0145-1)
Formic Acid, "FA" (Fluka cat # 55302-LC-GF)
Tris Buffer Saline, 10X, "TBS" final 1x, 50 mM Tris, pH 7.4, 150 mM NaCl (Becton BioProducts cat # BM-301)
HaltTM Proteases and Phosphatase inhibitor cocktail, 100X (Thermo ScientificTM cat # 78440)
Detection reagent: LuminataTM Forte western HRP Substrate (EMD Millipore cat # WBLV0500)

Antibodies (EMD Millipore):
Anti Alzheimer's precursor protein A4 (cat # MAB348)
Anti Tau phosphoSerine 396 (cat # AB9658)
Anti amyloid β clone WO-2 (cat # MABN10)
Goat anti mouse (cat # AP132P)
Anti Tau 4-repeat isoform RD4 (cat # 50-804)

Concentrators (EMD Millipore):
AmiconTM Ultra-4 (cat # UFC800308)
AmiconTM Ultra-0.5 (cat # UFC0308)

ELISA and MilliplexTM (EMD Millipore):

High sensitivity Human Amyloid β -40 ELISA 96 well kit (cat # EZHS40)
High sensitivity Human Amyloid β -42 ELISA 96 well kit (cat # EZHS42)

Human Amyloid Beta (6) & Tau magnetic bead panel 96 well plate assay (cat # HNABTMAG-68K)

Equipment and supplies

Acuity UPLC, coupled with XevoTM G2-S mass spectrometer (Waters)
Centrifuge with swinging buckets rotor (Beckman Coulter AllegroTM 25R; 15 mL tube adapter)

Direct DetectTM spectrometer (EMD Millipore cat # DDHW000-10-WW)

Trans-BlotTM SD-Semi-Dry Electrophoretic Transfer Cell (Bio-Rad cat # 170-3940)

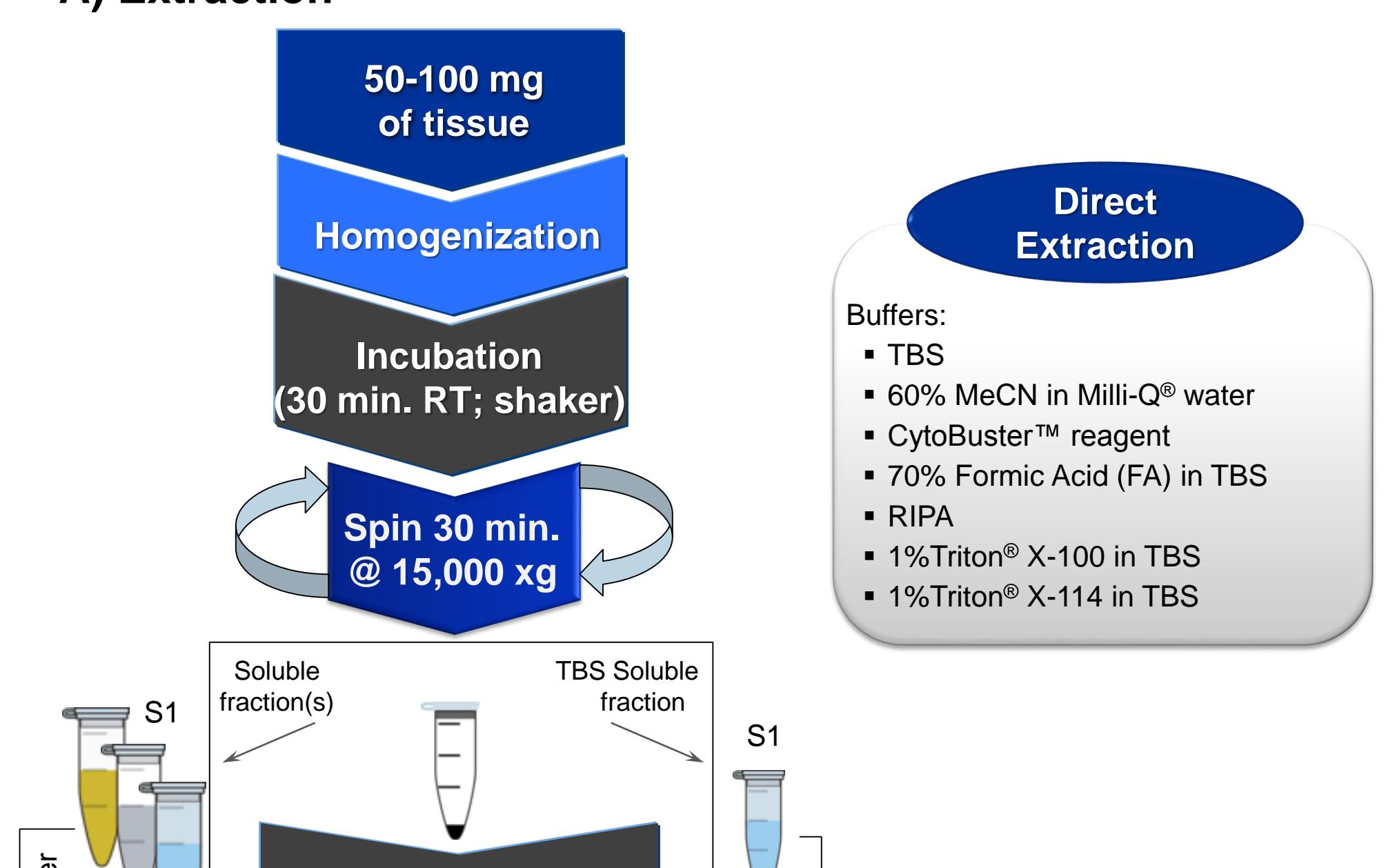
SNAP i.D.TM 2.0 Protein detection system (EMD Millipore cat # SNAP2BASE)

NuPageTM 4-12% SDS PAGE gels (Thermo ScientificTM)

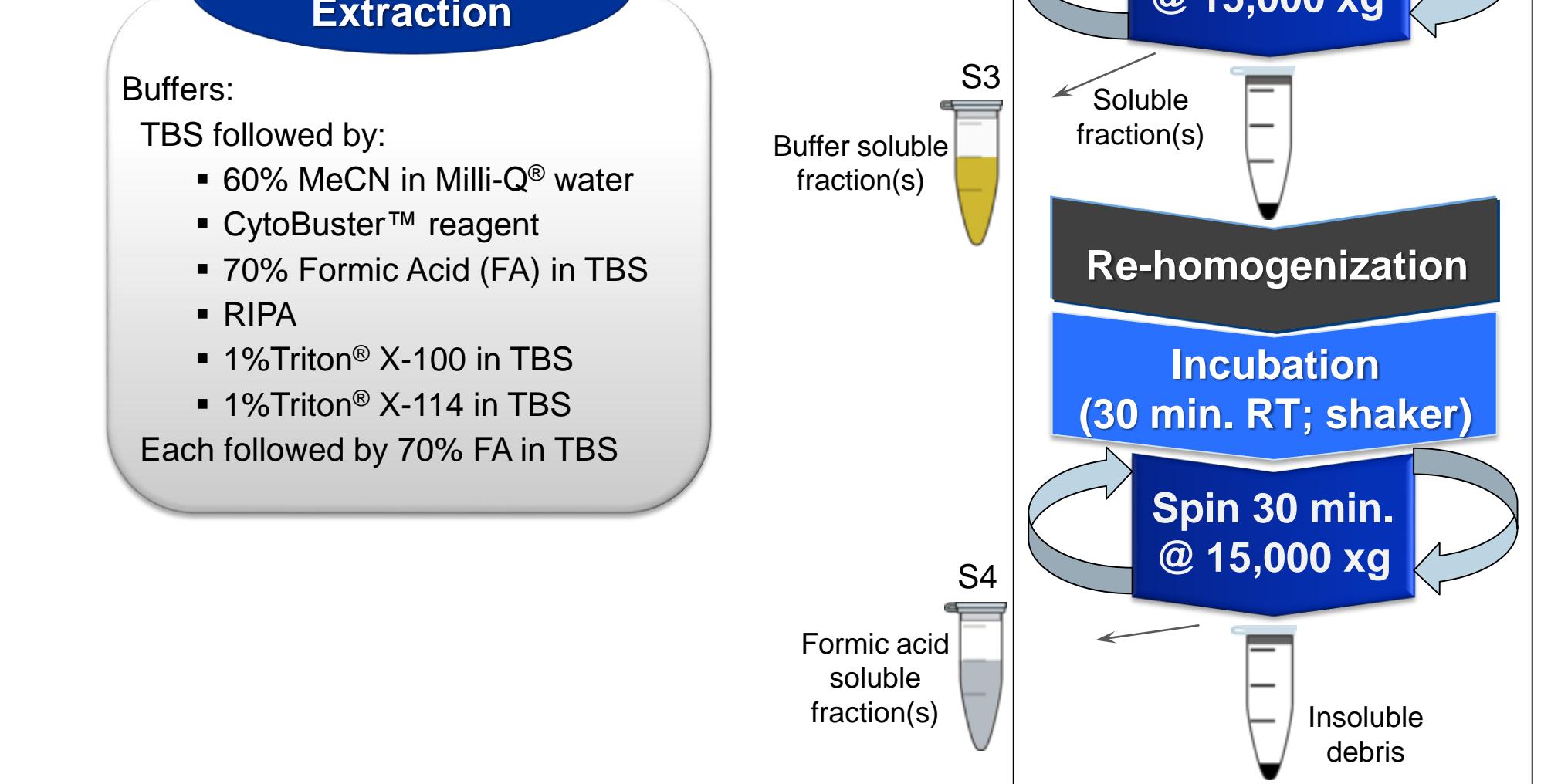
ImmobilionTM-P membrane (EMD Millipore cat # IPVH08130)

Methods

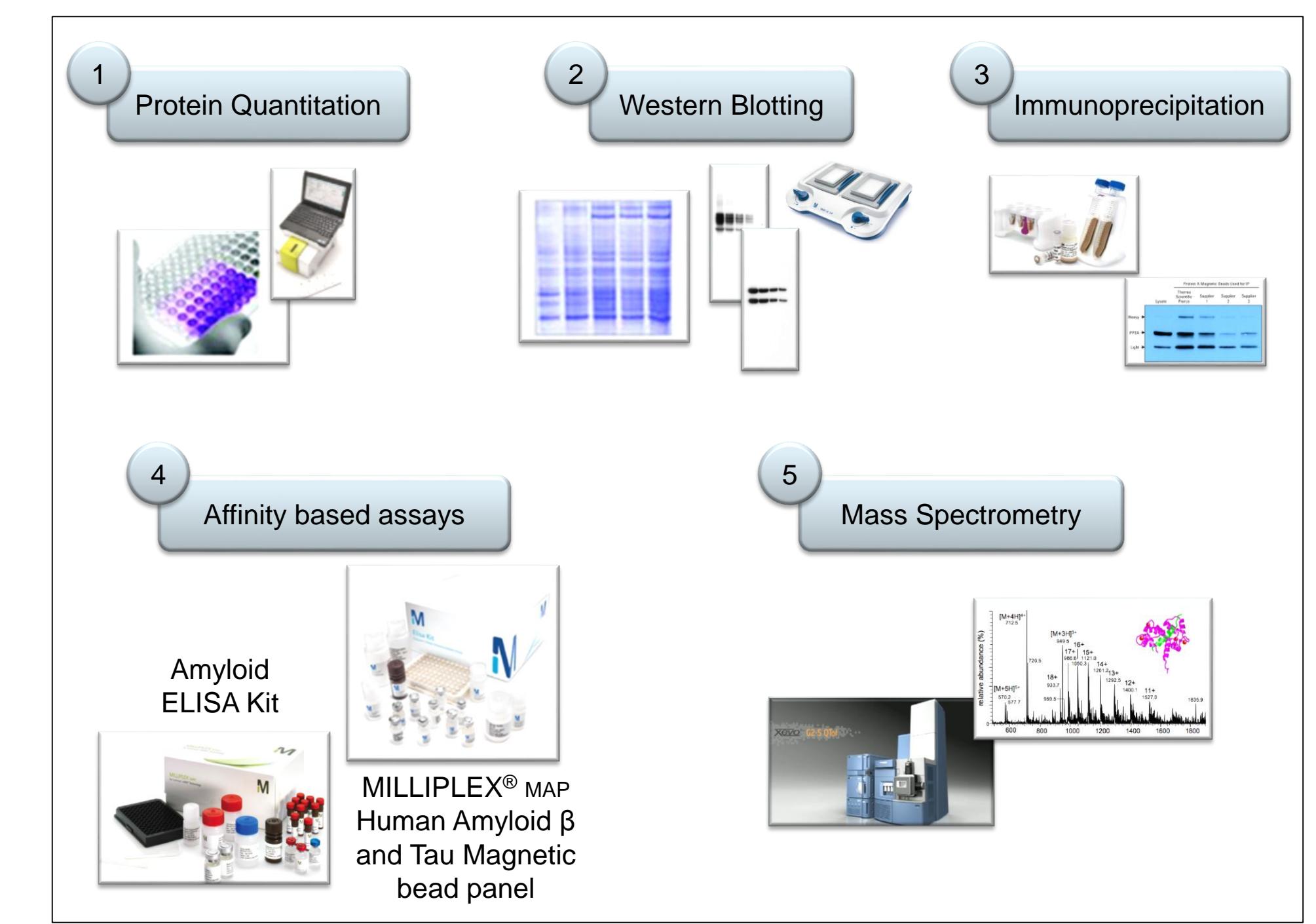
A) Extraction



Sequential Extraction



B) Analysis



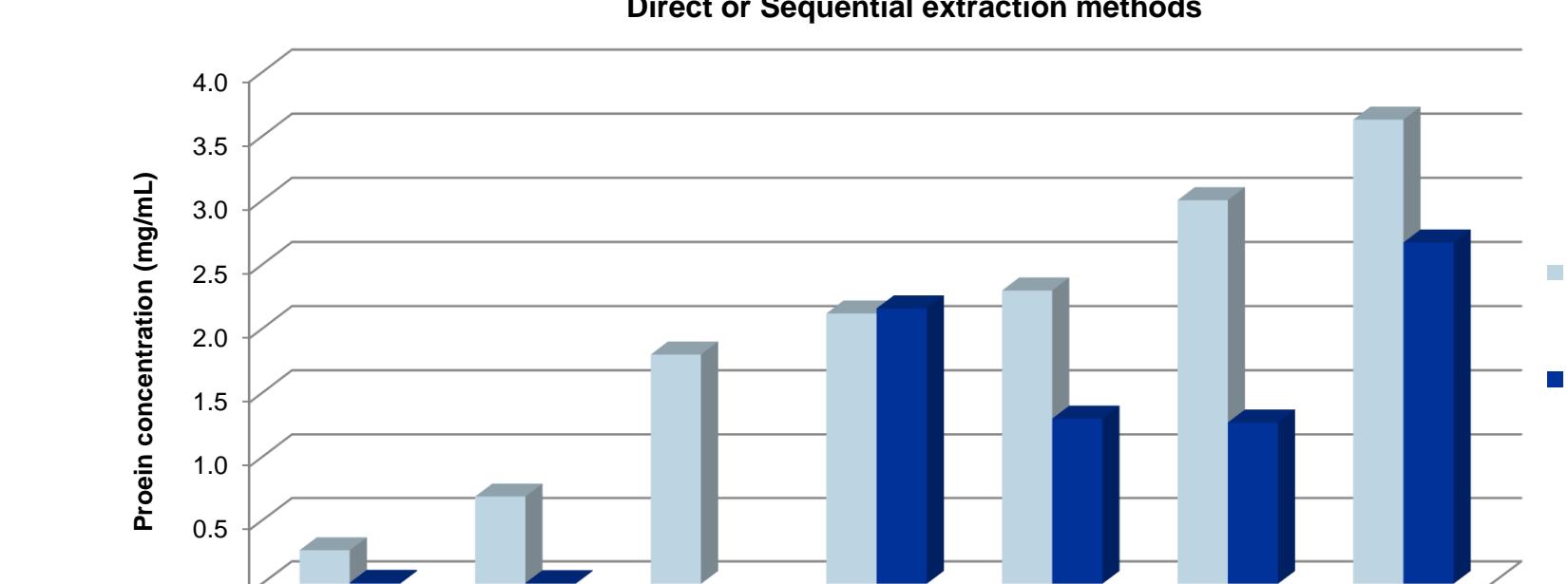
Results

Total protein concentration measured by Direct DetectTM spectrometer

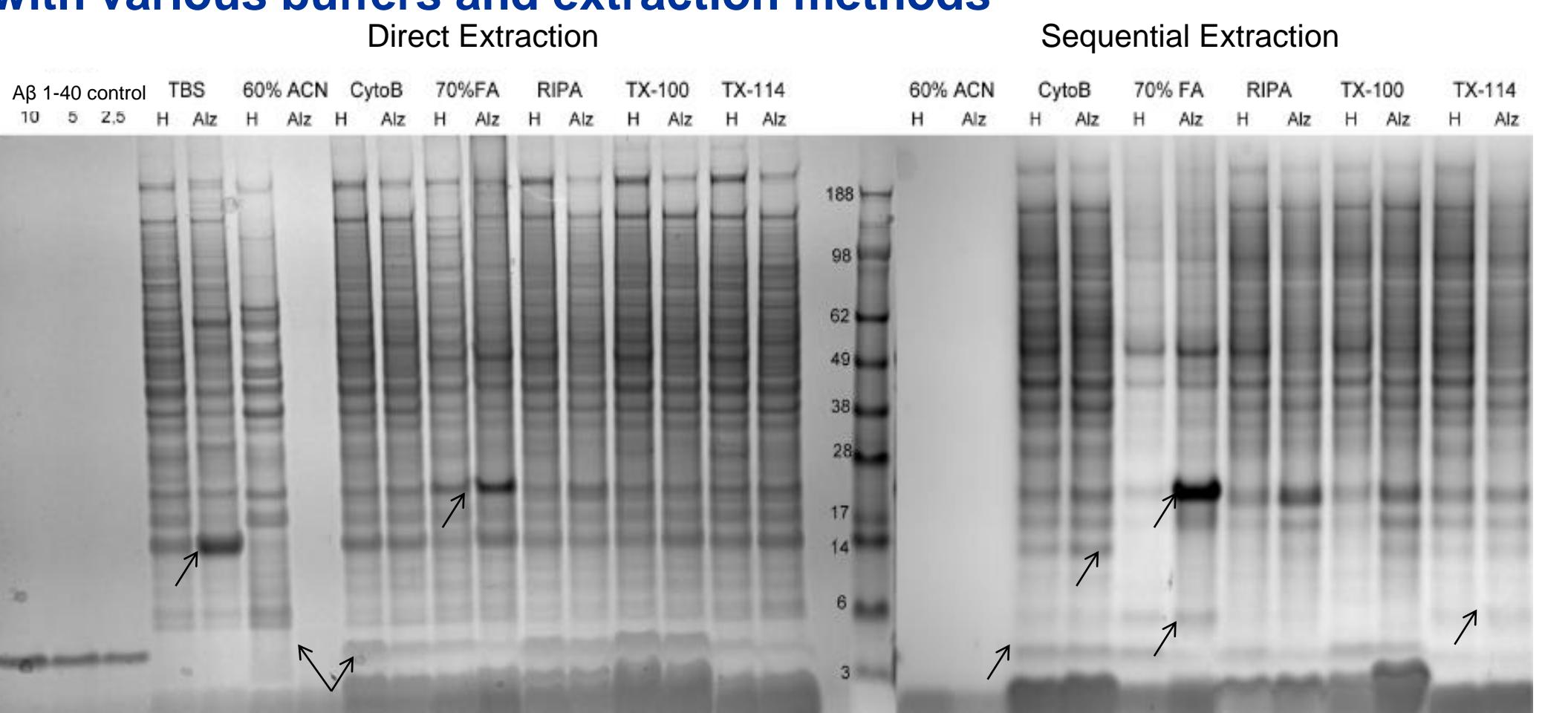
Analysis of total protein concentration in the lysates was used to assess efficiency of the applied extraction method. Results, presented below, show that application of a single detergent or a mixture of ionic detergents allows for efficient protein extraction in a single step (Direct Extraction).

	70% FA (mg/mL)	60% MeCN (mg/mL)	TBS (mg/mL)	1% TX-100 (mg/mL)	1% TX-114 (mg/mL)	CytoBuster TM Reagent (mg/mL)	RIPA (mg/mL)
Direct Extraction	0.3 ± 0.5	0.7 ± 0.9	1.8 ± 0.7	2.1 ± 0.2	2.3 ± 1	3.0 ± 0.3	3.6 ± 0.9
Sequential Extraction	Not detected	Not detected	Not Applicable	2.1 ± 1	1.3 ± 0.3	1.3 ± 0.3	2.7 ± 0.2

Total protein concentration in brain tissue lysates prepared with various buffers using Direct or Sequential extraction methods



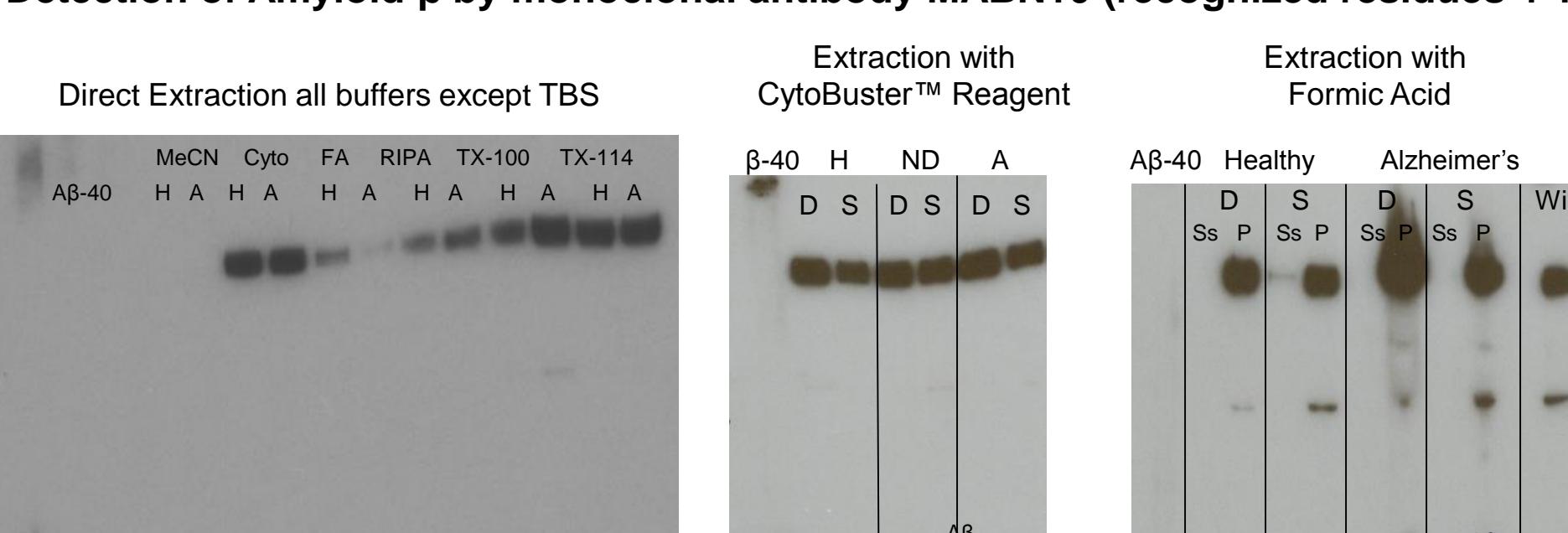
Gel electrophoresis of protein extracted from human brain tissue with various buffers and extraction methods



There were clear differences in protein profiles depending on buffer used for the extraction. The arrows point at some of the more significant differences between types of preparation.

Detection of neuronal markers by SNAP i.d.TM Western Blotting system

Detection of Amyloid β by monoclonal antibody MABN10 (recognized residues 4-10)



Alzheimer's precursor (APP) was detected in all lysates, however higher content was observed in samples prepared using the Direct (D) Extraction method.

H = Healthy
ND = Not Diagnosed
A = Alzheimer's

D = Direct Extraction
S = Sequential Extraction
Ss = Suspension
P = Precipitated fraction

Healthy ND Alzheimer's

D S D S D S D S

Aβ Aβ Aβ Aβ

Aβ Aβ Aβ Aβ