Institutt for kjemisk prosessteknologi NTNU

Poster presentation

Presentation of the work:

The main results of the project work are to be presented on a poster. Each group shall make a poster with the size 0.7×1 meter. The posters are to be presented in the corridor between the building 5 and the Hall (outside the auditorium K5).

Monday 22nd November at 1300-1500

At least one of the group members must be in the vicinity of their poster at any time to answer any questions related to the work.

Some practical information:

A short description on how to make a poster is attached. We have no access to a printer that is able to produce paper size 0.7×1.0 m format. Therefore you have to print out on A4 or A3 formats and then mount the sheets on a cardboard. You will get a cardboard from the department office. Since you have no access to a color printer, ask your advisor to print out color sheets if needed. Here, the poster should be in portait format, i.e. height 1m and width 0.7m. You should use portrait format, i.e. height is 1.0 m.

How to make a poster

On engineering and scientific conferences it is very common to use posters to communicate results. Compared to an oral presentation, a poster presentation may be an equally effective way of communicating. Since you are present at your stand, you will meet with the people that have interests in your work. A poster should catch the interest of these people in 3-4 seconds; otherwise they will pass your stand. The design of the poster is therefore very important. Even more important is the content, which should give the impression you have made an investigation of the problem on a relatively high level, and that the results are significant.

The poster will normally contain the following elements:

- Title
- Authors (with affiliation)
- Introduction
- Problem definition and confinement
- Method of investigation
- Results
- Conclusions and recommendations

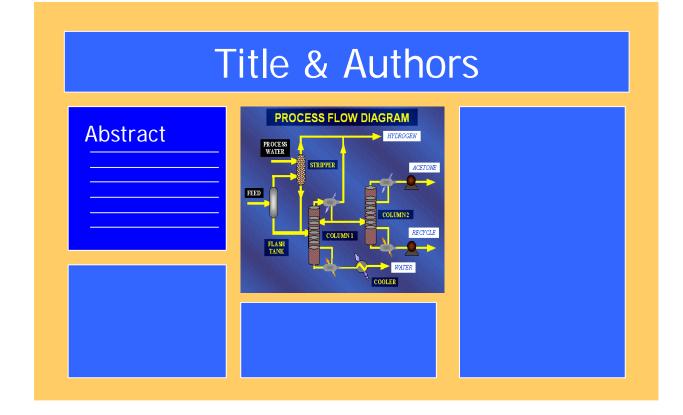
These items are not necessarily headlines, but rather elements that should be there. How much space you devote to each element will depend on the work that you are presenting.

Some hints:

- *Present the problem and results straight forward and simple.* There is not enough space for all details on a single poster. (However, you may bring with you a paper with extra details for those who are interested)
- *The text should be readable from a distance of 2-3 meters.* You should choose a font size and type that is easy to read. The font size of the main text should be minimum 25p, headlines 45-50p and the title about 100p.
- *You may use colors, but do not exaggerate.* Text and colors should have a good contrast to the background. Use complementary colors.
- *Figures, plots and process flow diagrams are important.* Axes must be labeled and have units. Try to simplify without loosing too much essential information, in particular process flow diagram (PFD) of large processes. The font size on figures should be on same size as the text.
- *The poster should attract the audience to your stand.* But remember that the conversation with the audience is even more important than the design of the poster.

Attached are two examples of how the poster layout might look like, and two scientific posters.

GOOD LUCK!



Title & Authors

| Problem | Methods | Results | Table 1 |
|---------|---------|---------|-------------|
| Goals | | | Fig 2 |
| | | | Conclusions |

Activation of p38 kinase in Atlantic salmon macrophages

Audny Johansen & Jorunn B. Jørgensen, Fiskeriforskning, Norwegian Institute of Fisheries and Aquaculture

MATERIALS AND METHODS

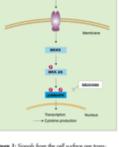
ends. After atimulation cells were lysed in SDS sample buller. The cell lysate was resolved by NuPAGE 10% Bis-Tris gel (Plaved), foi-lowed by transfer to PVD-membrane (Millipote). The filter was incu-bated with phosphaspecific DBS MAPK enthody lab) (PVE England Biolobu) and immunodetection was performed as instructed by the manufacture. The membranes were stripped and reprobed with a nabbit polyclonal ab against actin (Sigma) and detected. The Kodok Digital Science TM Image Station was used for making digital images of the membranes. Kodak Image Analysis Software was

The p38 kinase is a member of the Mitogen-activated protein kinase (MAPK) family of proteins. The MAPKs are highly conserved during evolution and are found in all eukaryotes from yeast to mammals (reviewed in 1). The p38 signalling transduction pathway plays an essential role in regulating many cellular processes including inflammation, cell differentiation, cell growth and death. Activation of p38, often through extracellular stimuli such as bacterial pathogens and cytakines, mediates signal transduction into the nucleus to turn on responsive genes (fig. 1). The p38 signalling pathway is shown to play an essential role in the production of inflammatory cytokines such as IL-13, TNF-a and IL-6 (reviewed in 2).

To know whether p38 is activated in Atlantic salmon macrophages, we used Western blot analysis with antibodies against the Thr180/Tyr182-phosphorylated form of p38 to detect its activity.

AIM

LPS, PMA, synthetic CpG oligodeoxynucleotides (ODNs) and synthetic doublestranded (ds) RNA poly (IC) were tested for their ability to activate p38 in salmon macrophages.



Stress, LPS

satory Cytol

Figure 1: Signah from the cell surface and duced through the cytoplasm by a cascode of pro-tein kineses: MAPK kinese kinese (MMOKL), MAPK kinese 3/6 MMOKC/4) and g3R MAPK. SB 203580 is a selective inhibitor of p3B kinase.

COURSE AND A

Materials

Phosphothi ate modified CpG ODNs were purcha used from Medprobe. Lipopolysaccharide (JPS), the synthetic ds RNA poly (IC) and phorbal myristate phosphate (PMA) were from Sigma. The inhibitor S8203580 was from Alexis. Western blotting

Atlantic salmon head kidney macrophages were cultivated for 3 days in L-15 medium with 5% FCS. Then medium were replaced with serumfree L-15 and stimulated as indicated in the figure leg-

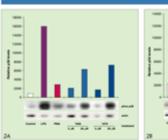
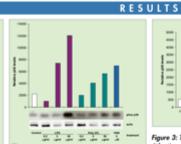


Figure 2A/B: CpG-DNA, LPS, poly (K) and PMA activate p38-kinase in Atlantic salmon macroph (2A) Atlantic so ophages were st lated for 2h with CoG ODNs 1668 and 1670 at two concentrations (5 and 20 µM), US (50 µg ml ⁻¹) and PMA (200 ng ml ⁻¹) or



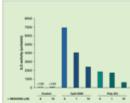
(28) LPS, poly (IC) and CpG ODN 1668. Controls are untreated cells. After stimulation

cells were lysed and subjected to Western blot analysis using antibodies described in M&M. Results shown are representatives of three separate experiments.

SUMMARY

Figure 3: Time course of p38 kinase activity in mocrophages stimulated with

LPS or poly (IC). Macrophages were stimulated with LPS or poly (IC) at 30 µg ml⁻¹ and harvested at indicated time after exposure. After stimulation cells were lysed and subjected to Western blot analysis using antibodies described in M&M.



used to analyse the images, to enable p38 level to be expressed rel-ative to actin expression in each sample.

Cell protection test for measurements or annual activity (MSE cells treated with supervision from simulated inucceytes, were infected with (IPNV (Sp) at multiplicity of infection [m.o.k.] at 0.01 and 72 hours post infection cytograftic effects (CPE) were measured in microtite assay (3), Results are presented as interferon like cytokine activity (ILC). One unit is defined as the resiprocal dilu-tion at which 50% protection against virus infection is obtained.

Cell protection test for measurement of antiviral activity

Figure 4: CpG ODN and poly (IC) induced nt on p38 activity

cytokine release is dependent on p38 activit Atlantic salmon leucocytes were preincubated for 1h with the indicated concentrations of the p38 inhibitor \$8203580 (4). Cells with or with out \$8203580 were then stimulated for 48h with CpG ODN (5 μM) or poly (IC) (10 μg ml ⁻¹). The supernatants were then harvested and assayed for ILC-activity. There was no direct effect of the inhibitor in the concentrations tested on CHSE-cells or the ability of the IPN virus to replicate in these cells (results not shown).

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 p38 kinase activity can be detected in Atlantic salmon macrophages by a phosphospecific p38 antibody.

· LPS was shown to be the most potent inducer of p38 activity (>15-fold induction), followed by CpG ODNs (>8-fold induction), poly (IC) (>2-fold induction) and PMA.

 p38 activation was detectable 30 min after stimulation, and there was a further increase in activation after 2h of stimulation.

· Preincubation of leucocytes with the specific p38 MAP kinase inhibitor, \$8203580, inhibited production of interferon like cytokines by CpG ODNs and poly (IC). This finding suggest that p38 MAPK activation participate in signalling pathways in Atlantic salmon leucocytes.

Hanner, C., Gilson, S., Jopa, M. & A.G. L. Advan-Kalaure, C. J. Han, (2000), The p.W. signal manufacture much: T, Tavity, C. & P. de Kolalin. (1997), Sparing n (1998). Singup Antonia Foreira Generation of a Forei Gener Models Anno Yoort to Univer. Physiological Review, 29, 1230-1224. Springer Antonian et al. Antonian. Glober pagande 23, 123. Springer Antonian et al. Antonian. Glober pagande 23, 124. Springer Antonian et al. Antonian. Glober pagande 24, 124.



Automatic paralogous gene detection and structural annotation of multigene families: application to the MYB family of transcription factors in *Arabidopsis thaliana*.



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Introduction

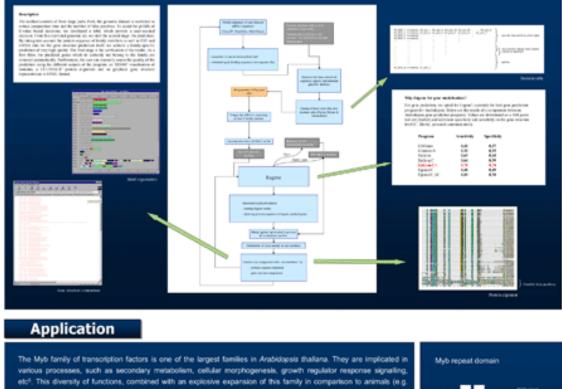
Now that more and more full genome sequences become available, the possibilities for evolutionary research seem endiess: one can collect all members of a gene family, without missing the low or conditionally expressed ones, which was often the case in cDNA-lbrary based family studies. Unfortunately, having the complete genome is only the beginning: the automatic annotation done by the large sequencing consortia is often of poor quality¹. As such, it is very unlikely that a perfect and exhaustive set of family members can be collected just by using the family name as query in a search engine (e.g. SRS). To avoid the tedious task of manually correcting structural and functional annotation, we developed a method that allows us to retrieve exhaustively all family members given a representative set (e.g. of experimental origin) and a set of genomic (e.g. BAC) sequences. The final result of this routine consists of the gene

structure, position and mRNA/protein sequence of all the family members.

This method was applied to the Myb family of transcription factors in the Arabidopsis thaliana genome.

Methodology

prediction was correct.



etc². This diversity of functions, combined with an explosive expansion of this family in comparison to animals (e.g. Humans have only 3 known Mybs), make this family very interesting for gene family evolution studies. As a family, it is not of the easiest to detect: the only conserved area is a 50as sequence which is repeated (degenerately) up to three times. At the genomic level, the repeats are usually interrupted by an intron, which increases the detection difficulty even further. From a set of 24 representative R2-R3 Myb subfamily members, ±140 R2-R3 Mybs were found in the Arabidopsis genome, which is in agreement with a recent study done on Arabidopsis transcription factors⁹. In more than 90% of

the cases, the comparison to the other family members allowed us to decide that the automated gene structure



Research Sciences

A DESCRIPTION OF A DESC

Bekromens, I. Auburg, S. and Rosse, P. (2001), Plane Physick, Biochem. JR, 1-13. L Group J. et al. (1997) Comput. Appl Biosci, 1368, 401-8. X. Theorynon, J.D. et al. (1997) Nucleic Acids Research 24, 4876-82. A. Schler, T. et al. FORM/2003 Charmin Convertes Biologic Informations (Multimotiones), T. et al. (2003). Science 13, 2003 Science 13, 20