



Validation of Genetic Determinants of Skeletal Diseases

Dr Kasia Piróg
(Newcastle University)



What is ?

Partners

Name

Newcastle University

University of L'Aquila

University of Manchester

Alacris Theranostics

University of Pavia

Polygene

Consiglio Nazionale delle Ricerche

Institut National de la Santé et de la Recherche Médicale

Certus Technology Associates Limited

Charité - Universitaetsmedizin Berlin

GATC Biotech AG

University Medical Center Hamburg Eppendorf

Evercyte (EVCYT)

University Hospital of Cologne

PRIMM Srl

University of Freiburg

University of Antwerp

Finovatis



Skeletal conditions

Common

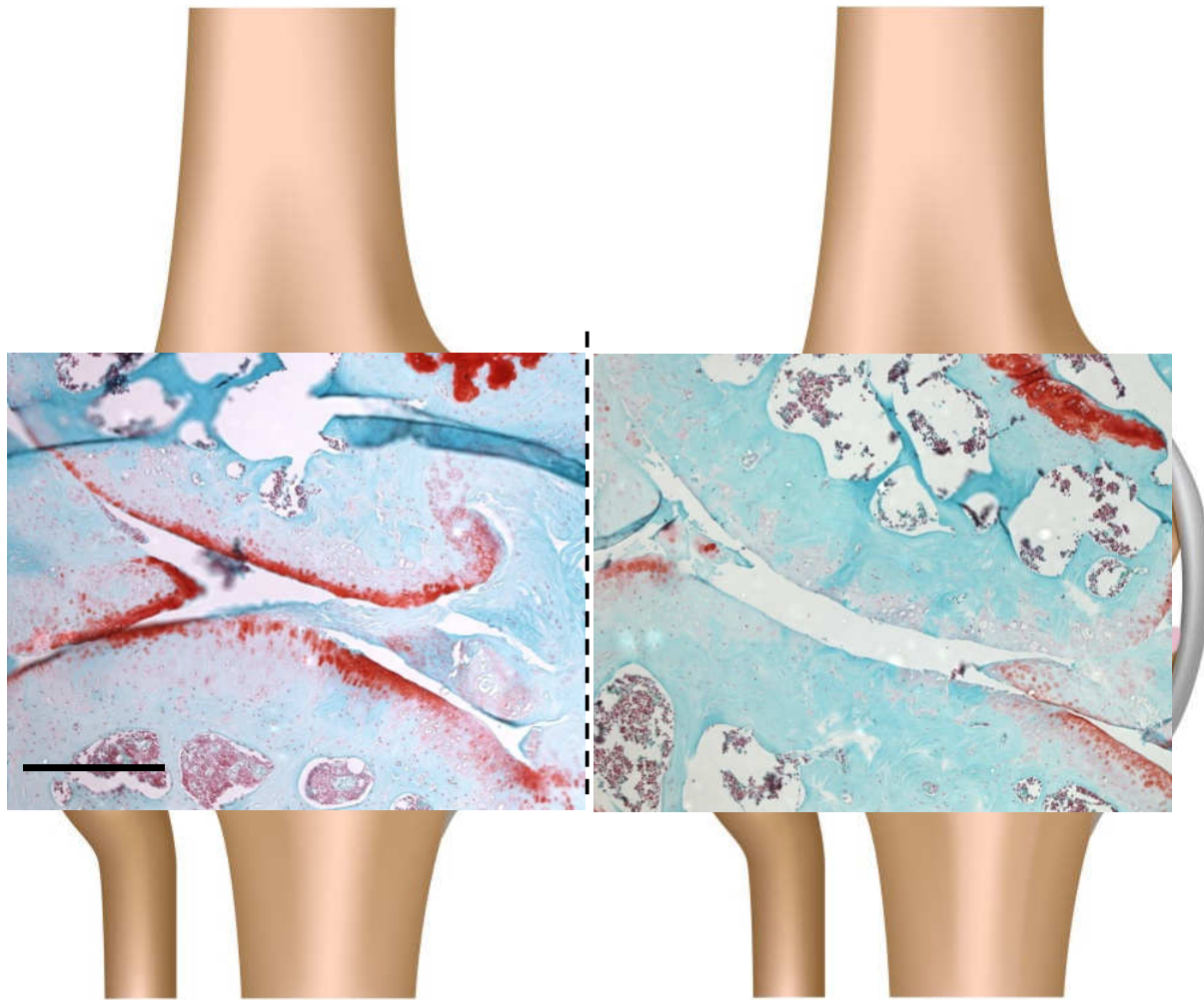
- related to ageing and/or injury
- often multifactorial
- often difficult to model
- SYBIL focuses specifically on osteoarthritis (OA) and osteoporosis (OP)
- **35-40 million people** suffer from **OA** in Europe
- the lifetime risk for OP fractures is 30-50% in women and 15-30% in men
- **OA and OP represent a major healthcare burden with the projected expansion of elderly population**



Rare

- **450** different rare skeletal conditions
- an overall incidence of **1/4000**
- extrapolates to **225,000 people** in the EU
- affect growth and development of the skeleton
- often monogenic (mutation in single gene)
- modelled in cells and in transgenic animals
- **a simple model to analyse the effects of genetic defects on disease progression**

Case study - osteoarthritis

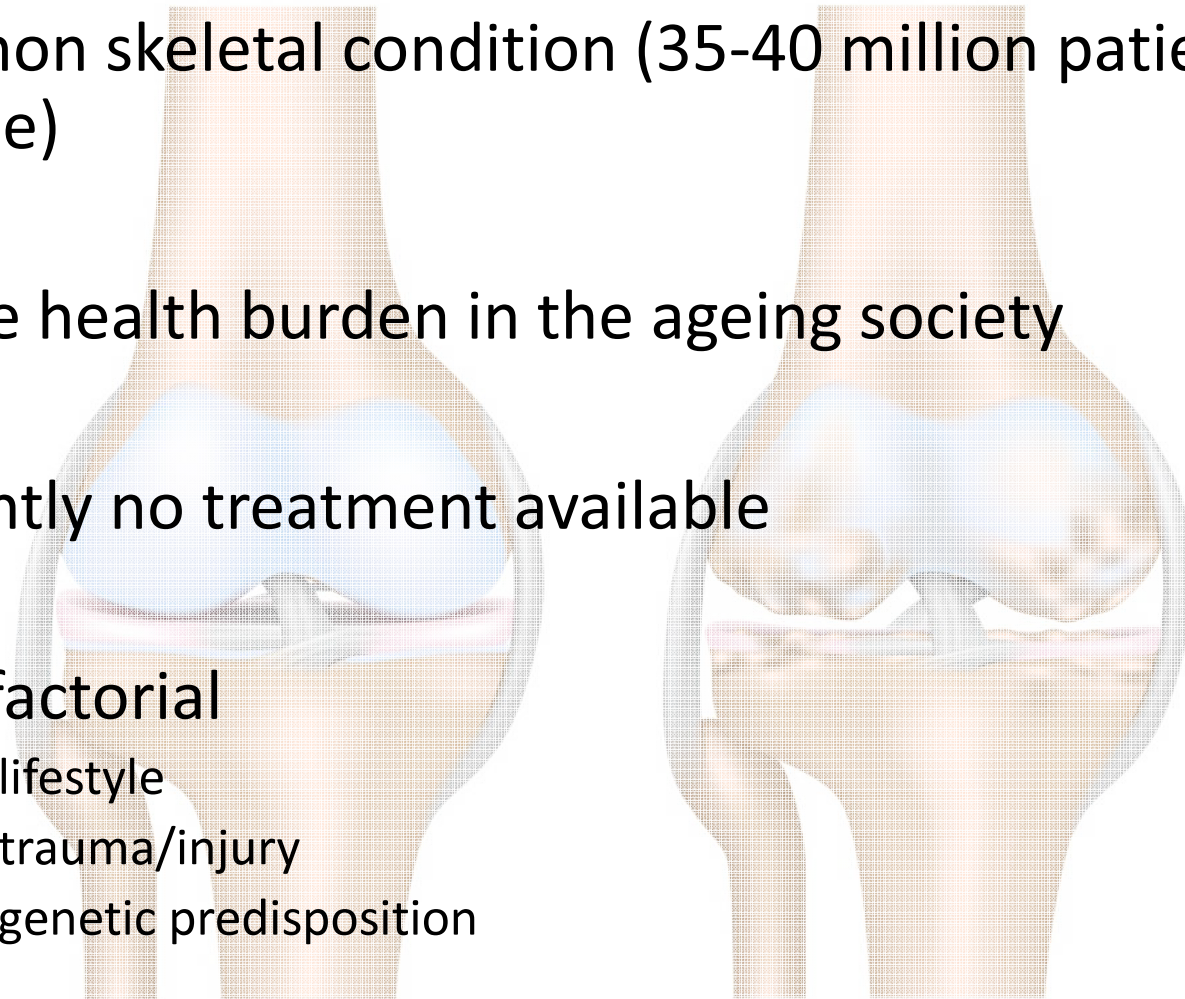


Healthy knee joint

Osteoarthritis

Case study - osteoarthritis

- common skeletal condition (35-40 million patients in Europe)
- severe health burden in the ageing society
- currently no treatment available
- multifactorial
 - lifestyle
 - trauma/injury
 - genetic predisposition
- modelled in animals often with surgical intervention



Rare skeletal conditions associated with osteoarthritis

- affect development and growth of the skeleton
- monogenic (mutation in a single gene)
- individually rare conditions but quite common as a group of disorders
- associated with musculoskeletal complications such as osteoarthritis
- easy to model in the laboratory setting

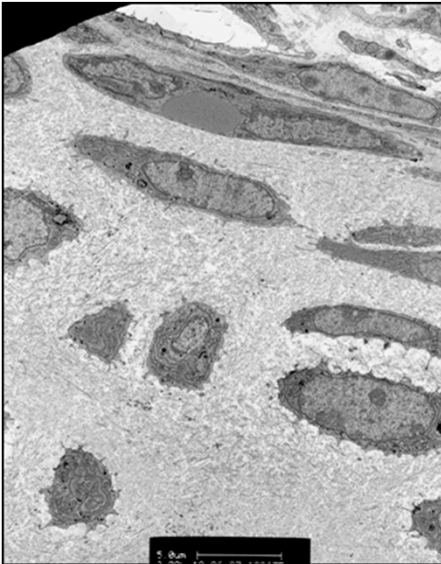
Multiple epiphyseal dysplasia



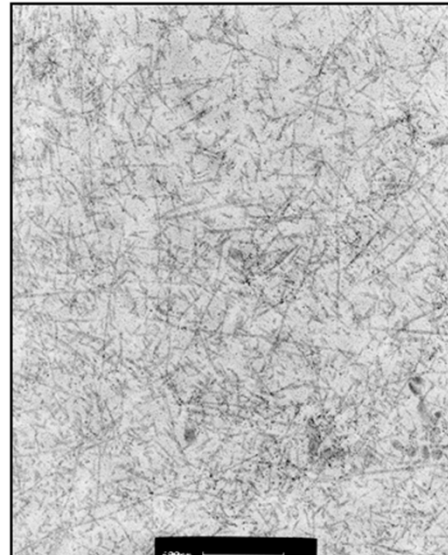
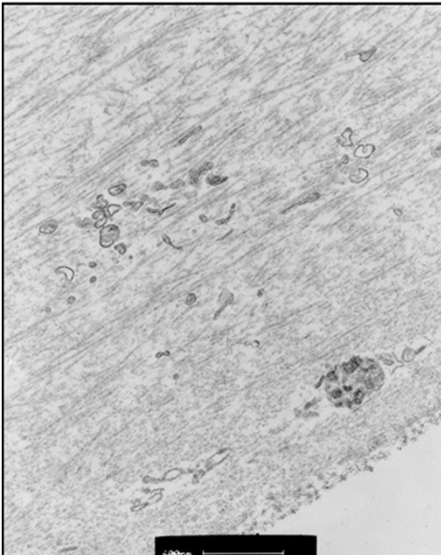
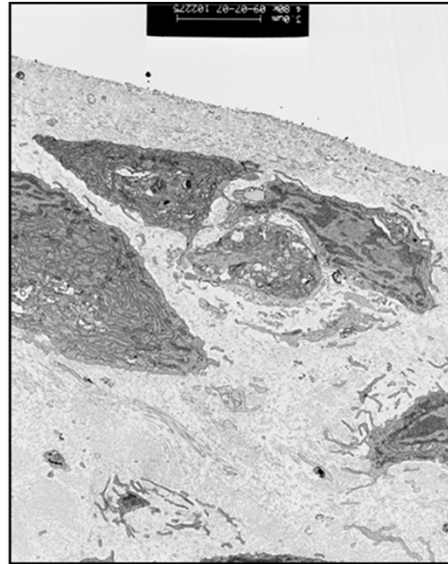
- autosomal dominant
- rare skeletal condition
- results from mutations in genes encoding structural proteins in cartilage
- short limbed dwarfism
- joint laxity
- **early onset osteoarthritis**

Mouse models allow the study of early degenerative changes

control

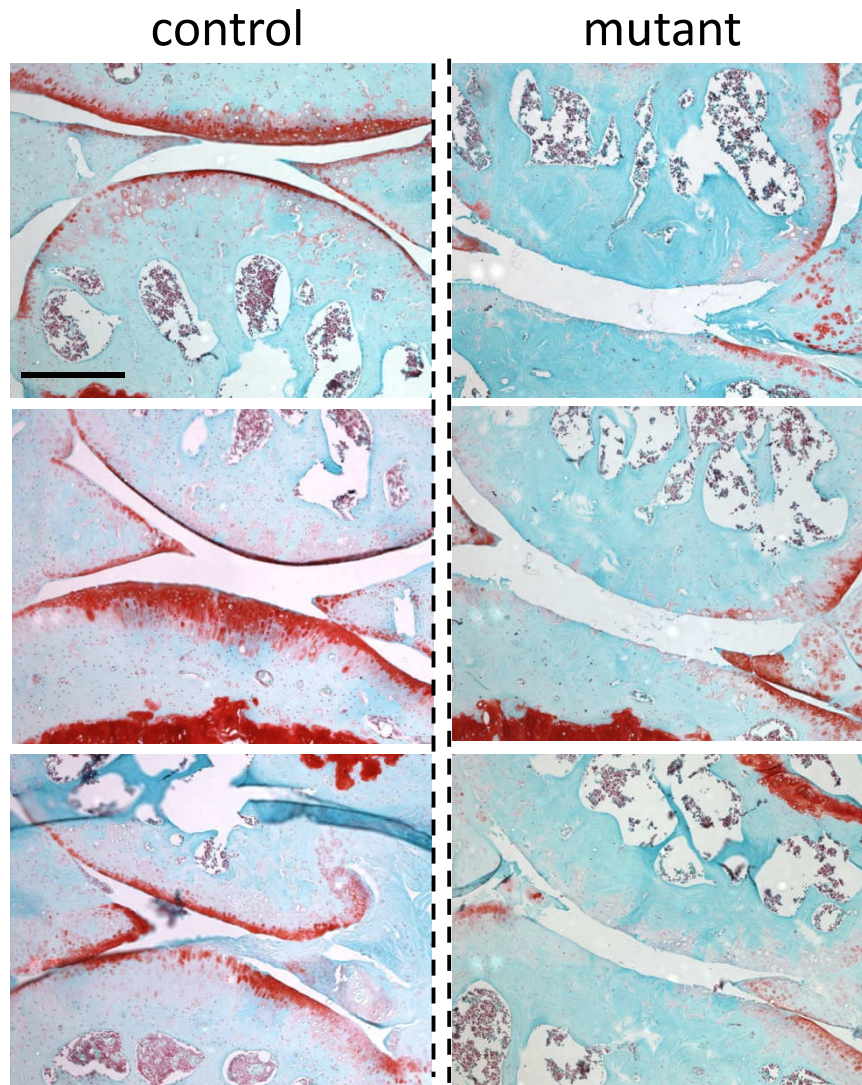


mutant



- first changes at 1 week
- abnormal cell death
- altered matrix composition
- altered cell morphology

MED mouse model develops early onset osteoarthritis



mouse

human

1 month

2.5 years

3 months

7.5 years

6 months

15 years

1 year

30 years

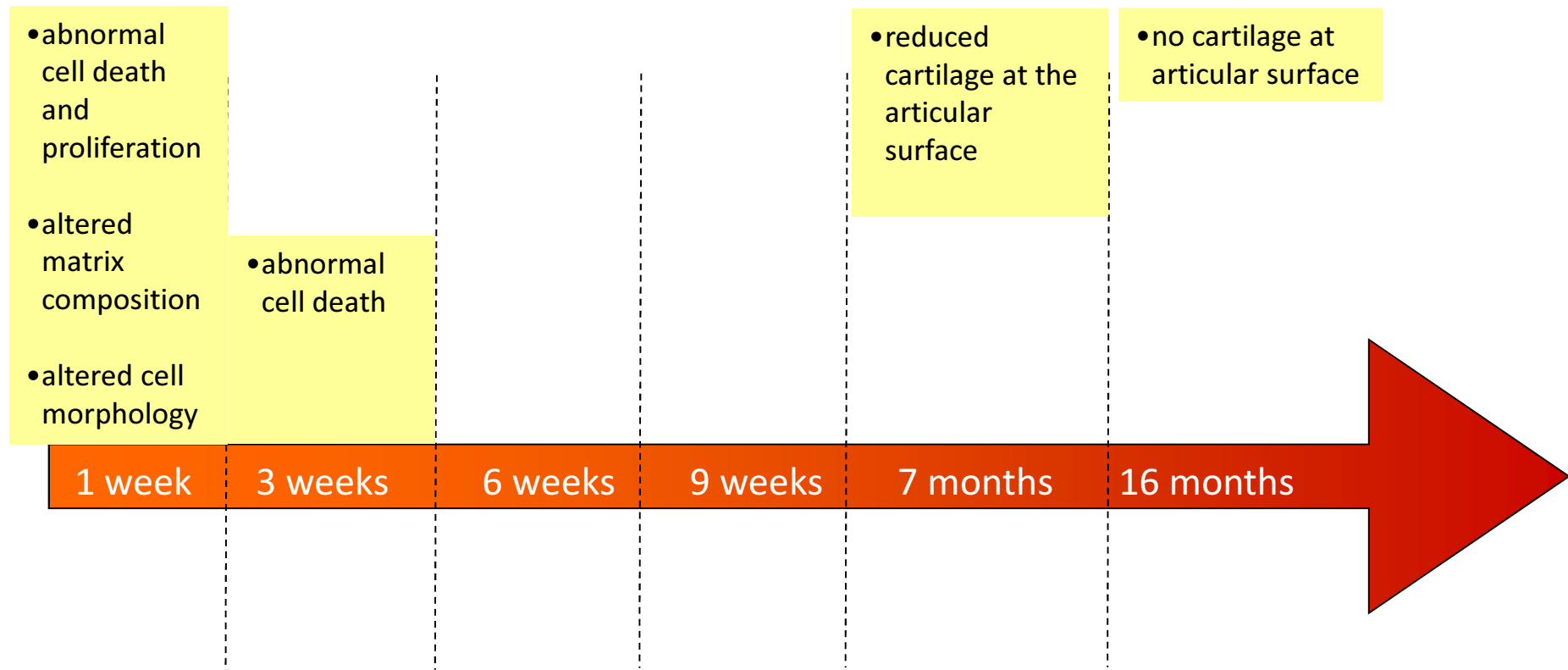
2 years

60 years

3 years

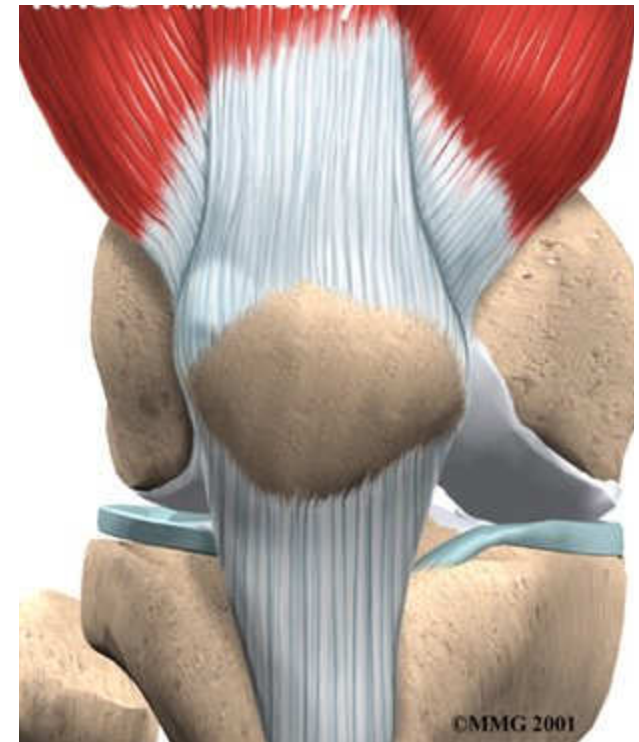
90 years

Mouse models allow the study of disease progression



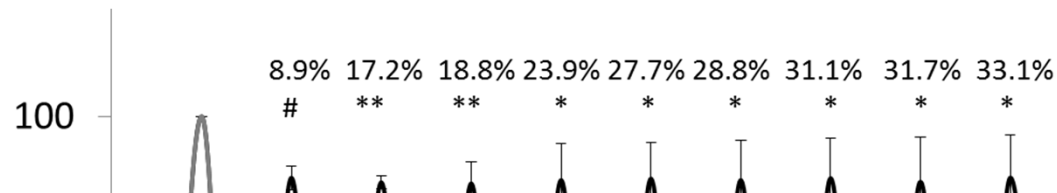
Soft tissue complications in osteoarthritis

- muscle weakness, tendon/ligament injury and joint laxity are the recognised conditions affecting the OA progression
- some MED patients are diagnosed with a “neuromuscular disorder” prior to correct skeletal diagnosis
- MED patients often suffer from tendon/ligamentous laxity

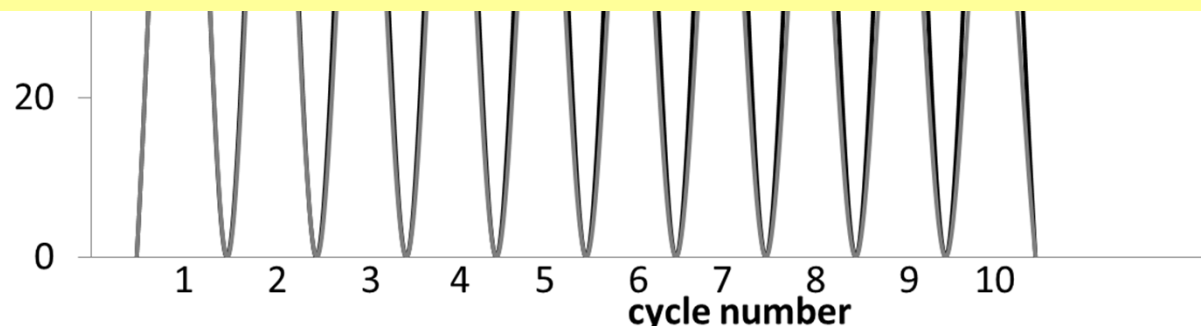


Soft tissue complications in osteoarthritis

Achilles tendons of mutant mice are more lax in cyclic testing



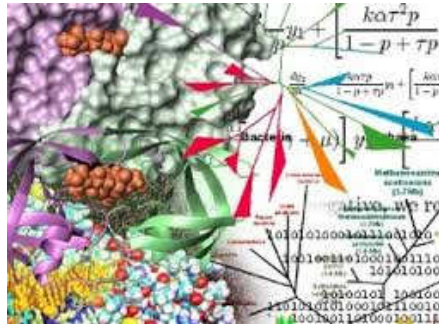
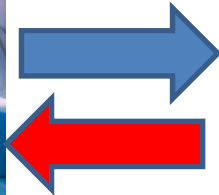
Mouse models of MED with structural changes in cartilage but no joint laxity do not develop early onset OA indicating a complexity in disease progression and providing a tool to dissect the disease mechanism in a relevant system



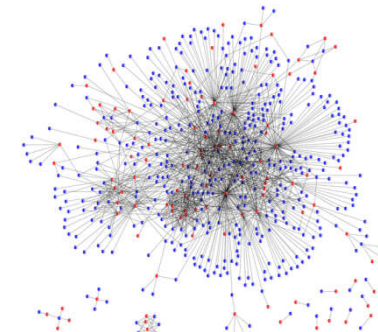
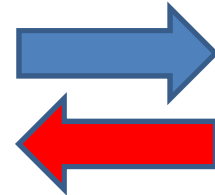
Systems biology approach



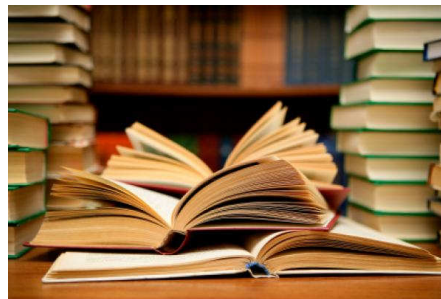
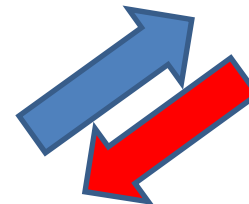
perform an experiment
*molecular behaviour in
bone/joint*



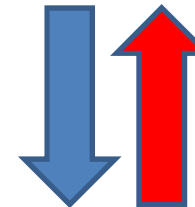
gather “omics” data
*gene expression,
proteomics etc.*



generate a network
*protein to protein or
gene interactions*

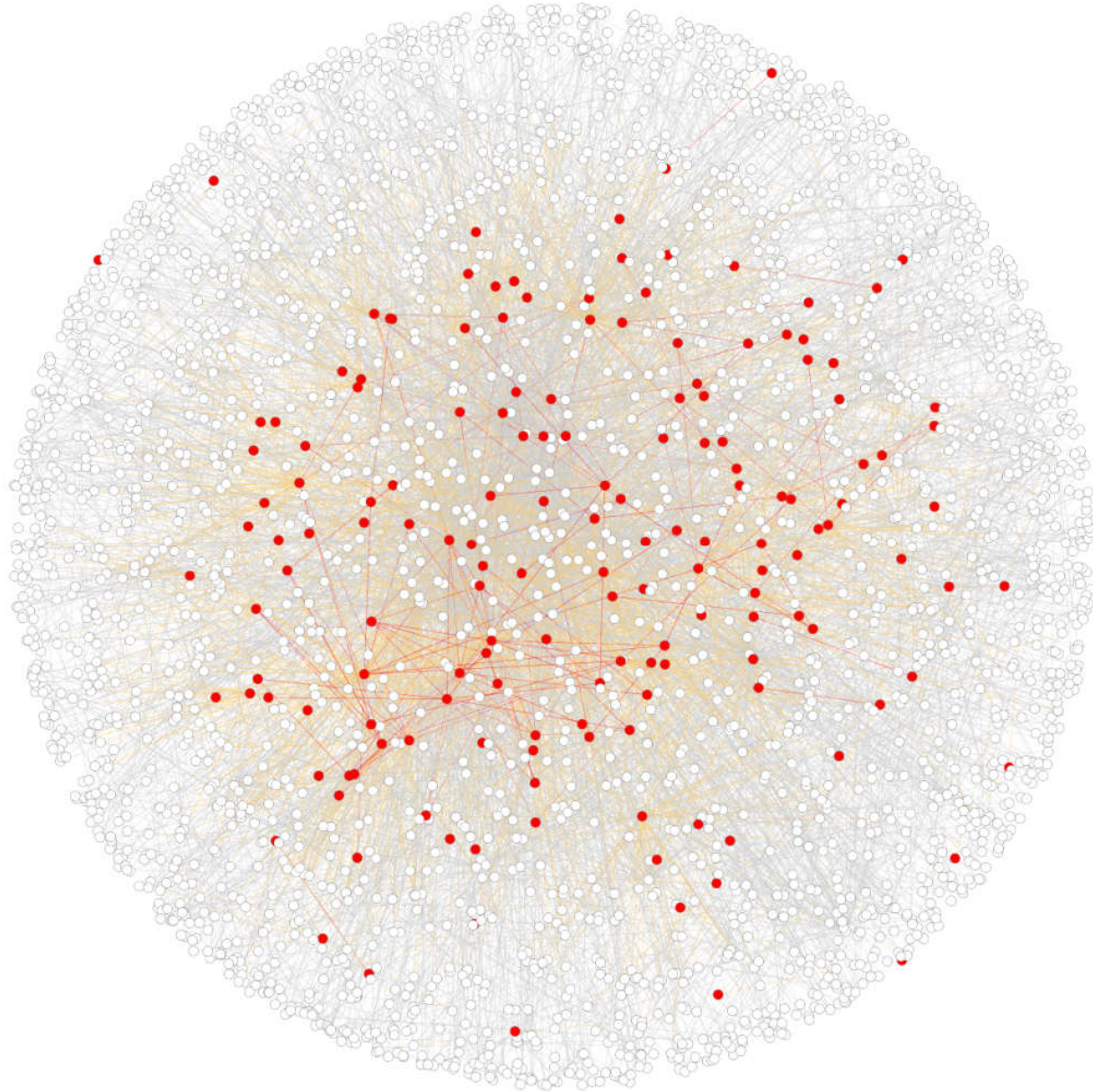


perform literature search
*genes of interest, “omics”
databases.*



delineate mechanisms
*diagnosis, treatment,
predictions*

Systems biology approach



What do we expect to achieve?

Functional validation of genetic determinants of skeletal conditions and ageing processes in animal and cellular models

- use animal and cellular models to study functions of genes associated with human diseases and/or ageing processes
- develop efficient, standardised and reliable tools and procedures
- identify, validate and create a portfolio of new biomarkers and therapeutic targets
- integrate and maintain the data in publically accessible web portals

For more information please visit:

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