

ICPerMed Best Practice in Personalised Medicine Award 2018. Berlin, 21 November 2018.



Navarra 1,000 Genomes Project (NAGEN 1000): *An example of a Project for Regional Implementation of Personalised Genomic Medicine in Healthcare*



Project with a nomination by ICPERMED AWARD
2018 "Best Practice in Personalised Medicine"

Angel Alonso
NAGEN Project Director.



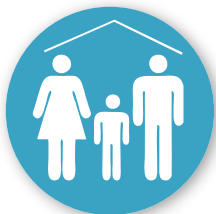
✚ "Best Practice in Personalised Medicine" Award 2018



Context



Personalised Genomic Medicine applications:



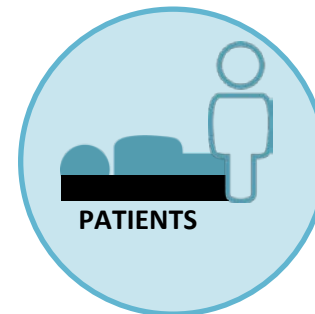
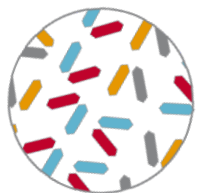
- > Diagnosis of diseases of genetic cause .
 - > Prediction, prevention
 - > Reproductive options
- > Characterization of genetic variants related to personal risk.
- > Identification carriers of serious genetic diseases.
- > Identification of genetic variants responsible for pharmacological response.
- > Stratification of patients for care management
- > Promote research and innovation.
- > Improve the quality of health care, better quality of life and save costs.

Aims NAGEN 1000



“Implementation of the use of information derived from WGS as a clinical tool for the development of personalised medicine in Navarre Public Health Service (SNS)”

- Translational research, innovation & genomics industry development



Sequencing 1,000 whole genomes from SNS patients with rare diseases.

Methodology



► Implementation Science Approach

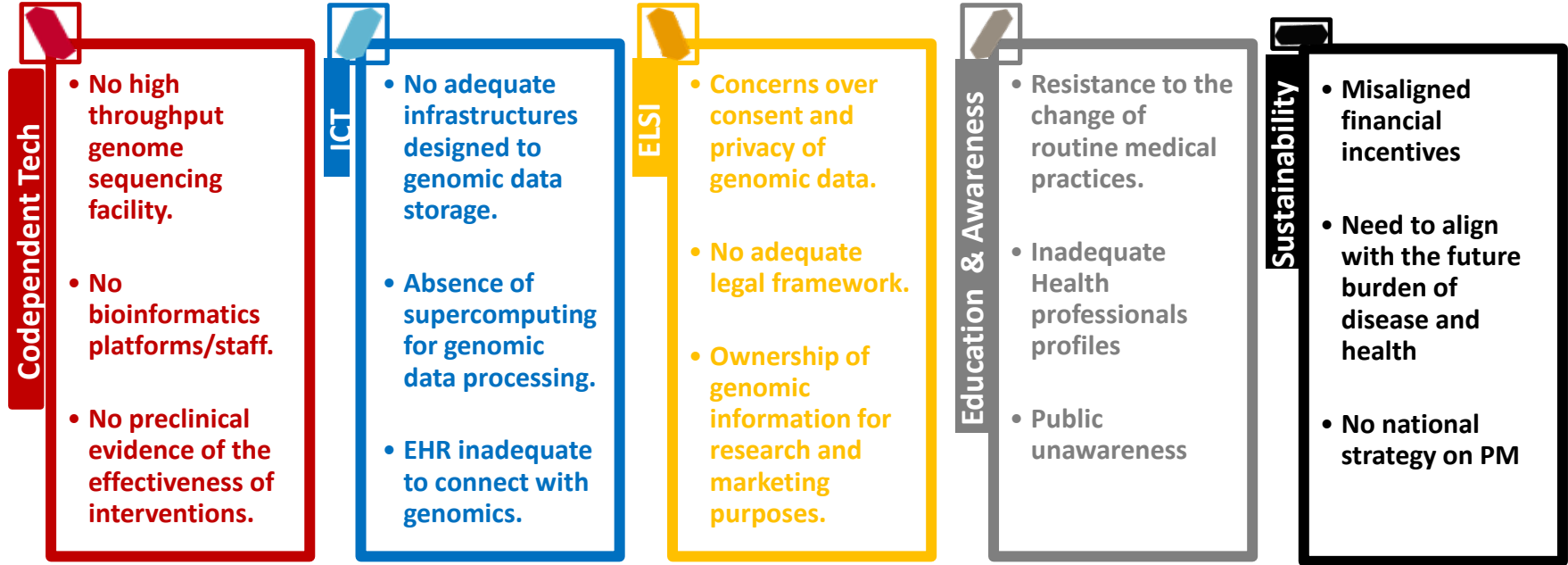


- > "It takes 17 years on average to convert the results of the research to benefit the patient"
David Chambers, director Implementation science National Cancer Institute.
- > "It is the method to promote the integration of research findings in health care" (NIH, 2016).
- > Tools: Pilot projects and Identification of Barriers.

Results: Barriers



Local barriers for Genomic Medicine Implementation in Navarra (NAGEN 1000 project):

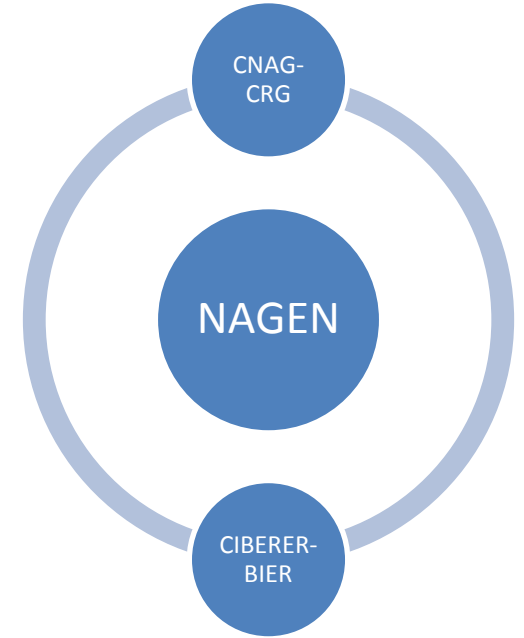


Results: Key Actions for this implementation



> Optimized use of pre-existing public infrastructures

- **CNAG-CRG** is a non-profit SEQUENCING PLATFORM funded by the Spanish Ministry of Science, Innovation and Universities and the Catalan Government .
- **CIBERER-BIER**, is a transversal BIOINFORMATICS PLATFORM funded by the Spanish Ministry of Science, Innovation and Universities and the Andalusia Government.





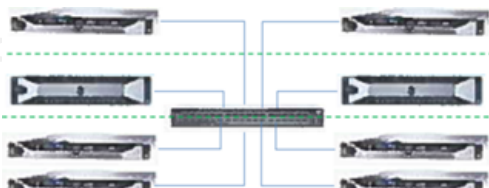
Results: Key Actions for this implementation

> > ICT

- 4 calculating nodes (IBM POWER 9 superprocessor).
- Infiniband transmission 100Gb/s

Optimizer

> Pre-clin



2x Managing nodes

2x HC Storage

4x Calculating nodes

Historia clínica Aplicaciones externas Fichas Usuario Listados Utilidades Ver Ayuda

NAGEN Entorno Hospitalario de Pamplona

Antecedentes Historia general Radiología Anál. Papan. Laboratorio Exploraciones Estimación

PROYECTO GENOMA - PROYECTO GENOMA DE INVEST. BIOP
EXPLORACION FUNCIONAL ALERGOLOGIA - 06/06/2017 - ALI
RESULTADO PROYECTO GENOMA NAGEN - 31/07/2018 - OTIC

Fecha Realiz.: F. Modif.: Caso: 0
Responsable: Servicio: NAGEN
Usuario soci.: Serv. Peri.: Serv. Peri.:

RESULTADO PROYECTO GENOMA NAGEN

Resultado 1 Se ha identificado la variante c.2477G>A, p.Ser225Asn en el gen WFS1 en heterocigosis, previamente asociado con el síndrome OMIM 602060 "Síndrome no sindrómico tipo 6, que es "muy probablemente" candidata a explicar el motivo de reclutamiento

RESULTADOS SECUNDARIOS PREDISPOSICION INDIVIDUAL A ENFERMEDAD

Técnica 2: Análisis bioinformático mediante panel virtual de 59 genes de predisposición a enfermedades genéticas propuesto por ACMG 2015

Resultado 2 No se han identificado alteraciones genéticas individuales con significado clínicamente relevante asociadas a predisposición a enfermedad en los genes analizados

RESULTADOS SECUNDARIOS DE RIESGO REPRODUCTIVO INDIVIDUAL

Técnica 3: Análisis bioinformático mediante panel virtual de los genes CFTR, FMR1 y SMN1 de riesgo reproductivo individual

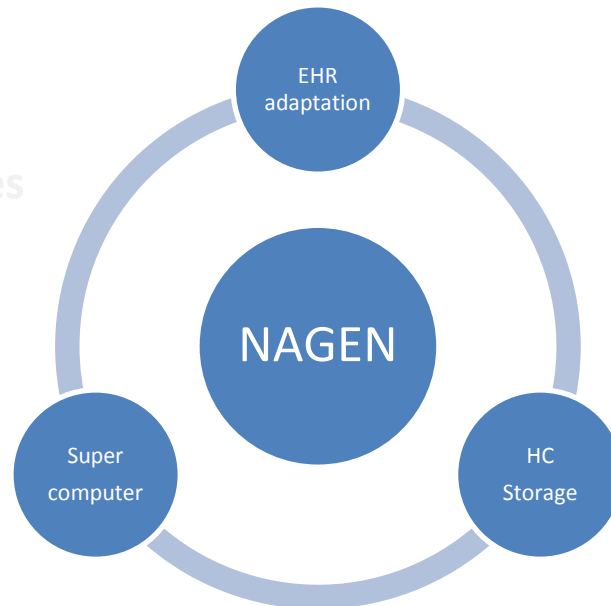
Resultado 3: No se han identificado alteraciones genéticas con significado clínicamente relevante asociadas a riesgo reproductivo individual en los genes analizados

RESULTADOS SECUNDARIOS DE RIESGO FARMACOGENÉTICOS

Técnica 4 Análisis bioinformático mediante panel virtual de los genes relacionados con una acción farmacológica condicionada por variación genética, según recomendación de la Sociedad Americana de Farmacogenómica, PhRMA/GfB)

Resultado 1 Medicamento: ATAZANAVIR (VH); dBSP: m837E29; Gen: UGT1A1; Variante en heterocigosis; Nivel de evidencia: (POKB) 1A; Genotipo: UGT1A1*1/1*10; Clasificación: toxicidad; Modificación: El tratamiento con atazanavir en estos pacientes presenta un menor riesgo de hiperbilirrubinemia que en la población general. La aparición de hiperbilirrubinemia puede depender de otros factores. Implicación en la prescripción: Principal

Alonso Sánchez, Ángel | NAGEN | 5384 | 1732





Results: Key Actions for this implementation

- > Education and training
- > Dissemination

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NAGEN monographic symposiumS (CHN, NB)	2
Hospital Clinical sessions CHN	1
Interhospitalary presentations (Comarcal, Primary)	3
Services clinical sessions	9
National Scientific Communications (CIBER,CNAG)	3
International Scientific Communications (British Society Genetics)	1
National Strategic Communications (Spanish Senate, Roche Institute, Sapanish Hospital CEOs meeting)	2
International Strategic Communications (DG Sante, EC)	1
International Concerted actions(CEIN, Pirepred)	2
General Public information (National Science Week, RD Patient association)	2
Press conferences and media difusion (news papers, radio, TV)	1
Website	1
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<https://www.nagen1000navarra.es/>

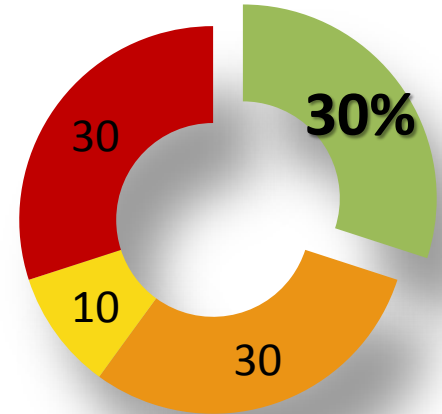


Results: Key Actions for this implementation



> Preclinical Evidence

Ref	Referral Code	Gen	Tránscrip	Heterozygosity	Mutation HGVS	OMIM	ClinVar
E0036.01	Hearing Loss	WFS1	ENST00000226760	Heterozygous	c.2477G>A p.Ser826Asn	600965 DEAFNESS, AUTOSOMAL DOMINANT 6; DFNA6.	
E0001.01	Balanced translocations	FAM227B	ENST00000299338	Heterozygous	46,X,Y,inv(15)(q21q24)		
E0009.01	Intellectual disability	KMT2A	ENSG00000118058	Heterozygous	c.654delA p.Ile218fs	605130. WIEDEMANN-STEINER SYNDROME	
E0009.01	Intellectual disability	KMT2A	ENSG00000118058	Heterozygous	c.656A>G p.Glu219Gly	605130. WIEDEMANN-STEINER SYNDROME	
E0015.01	Neurofibromatosis 1	SPRED1	ENST00000299084	Heterozygous	c.46C>T p.Arg16*	611431 LEGUIS SYNDROME	372718
E0017.01	Intellectual disability	PPM1D	ENST00000305921	Heterozygous	c.1274_1277dupGAGG, p.Asp425fs	617450 INTELLECTUAL DEVELOPMENTAL DISORDER WITH GASTROINTESTINAL DIFFICULTIES AND HIGH PAIN THRESHOLD; IDDGIP	
E0018.01	Hereditary Ataxia	LMNB1	ENST00000261366	Heterozygous	c.626A>G p.Lys209Arg	169500 LEUKODYSTROPHY, DEMYELINATING, ADULT-ONSET, AUTOSOMAL DOMINANT; ADLD	
E0024.01	Balanced translocations	FAAH2	ENST00000374900	Heterozygous	46,X,t(X;10)(p11,q26)	300654. FATTY ACID AMIDE HYDROLASE 2; FAAH2	
E0024.01	Balanced translocations	NHLRC2	ENST00000369301	Heterozygous	46,X,t(X;10)(p11,q26)		
E0026.01	Intellectual disability	WDR45	ENST00000356463	Heterozygous	c.971delT p.Val324fs	300894 NEURODEGENERATION WITH HIGH IRON ACCUMULATION 5; NBIAS.	
E0029.01	Familial Pulmonary Fibrosis	TERT	ENST00000310581	Heterozygous	c.1403G>C p.Cys468Ser	614742. PULMONARY FIBROSIS AND/OR BONE MARROW FAILURE, TELOMERE-RELATED, 1; PFBMT1	
E0001.01	Balanced translocations	SIN3A	ENST00000394947	Heterozygous	46,X,Y,inv(15)(q21q24)		
E0035.01	Undiagnosed metabolic diseases	ALDOB	ENST00000374855	Homozygous	c.448G>C p.Ala150Pro	229600 FRUCTOSE INTOLERANCE, HEREDITARY.	464
E0097.01	Kidney Proteinuria	INF2	ENST00000330634	Heterozygous	c.641G>A p.Arg214His	613237 FOCAL SEGMENTAL GLOMERULOSCLEROSIS 5; FSG5S	1053
E0040.01	Familial Breast Cancer	BRCA1	ENST00000357654	Heterozygous	c.5123C>A p.Ala1708Glu	604370, BREAST-OVARIAN CANCER, FAMILIAL 1 113705, BRCA1	55407
E0059.01	Artrorropisis	CRLF1	ENST00000392386	Heterozygous	c.713dupC p.Pro239Alafs	272430 CRISPONI/COLD-INDUCED SWEATING SYNDROME 1; CISS1	216913
E0059.01	Artrorropisis	CRLF1	ENST00000392386	Heterozygous	c.803T>C p.Phe268Ser	272430 CRISPONI/COLD-INDUCED SWEATING SYNDROME 1; CISS1	
E0059.01	Artrorropisis	CRLF1	ENST00000392386	Heterozygous	c.1018C>T p.Arg340Cys	272430 CRISPONI/COLD-INDUCED SWEATING SYNDROME 1; CISS1	
E0066.01	Noonan syndrome	PTPN11	ENST00000351677	Heterozygous	c.417G>C p.Glu139Asp	163950 NOONAN SYNDROME 1; NS1 and 156250 METACHONDROMATOSIS; METCDS	40513
E0077.01	Charcot-Marie-Tooth disease	SBF2	ENST00000256190	Homozygous	c.1281T>A p.Cys427Ter	604563 CHARCOT-MARIE-TOOTH TYPE 4B2; CMT4B2	
E0079.01	Charcot-Marie-Tooth disease	RAB7A	ENST00000265062	Heterozygous	c.484G>A p.Val162Met	602298. RAS-ASSOCIATED PROTEIN 7; RAB7; RAB7A. <i>Pathogenic Gene Symbol: RAB7A</i>	7346
E0080.01	Charcot-Marie-Tooth disease	MPZ	ENST00000533357	Heterozygous	c.148T>C p.Cys50Arg	159440. CHARCOT-MARIE-TOOTH DISEASE CMT1B, CMT2J, CMT2J, DEJERINE-SOTTAS, CHN, ROUSSY-LEVY DYSTASIA	
E0086.01	Epileptic Encephalopathy	SCN1A	ENST00000303395	Heterozygous	c.2585G>A p.Arg862Gln	607208 EPILEPTIC ENCEPHALOPATHY, EARLY INFANTILE, 6; EIEE6	
E0089.01	Familial hematuria	COL4A4	ENST00000396625	Heterozygous	c.735>2T>C	120131; COL4A4; Alport syndrome.	
E0093.01	Complex Parkinsonism	LRKK2	ENST00000298910	Heterozygous	c.4321C>G p.Arg1441Gly	609007; LRKK2, PARK8; Parkinson disease 8	1936
E0030.01	IUGR y IGF Abnormalities	IGF1R	ENST00000268035	Heterozygous	c.307C>T p.Leu103Phe	OMIM:270450 INSULIN-LIKE GROWTH FACTOR I, RESISTANCE TO, DUE TO INCREASED BINDING PROTEIN	



- Diagnosis achieved
- Strong candidate diagnosis
- Mild candidate diagnosis
- No diagnosis

Results: Key Actions for this implementation



> Preclinical Evidence

Secondary Results

Risk	Consent	Number of patients	% of cases
Disease risk	211	11	5,2%
Reproductive Risk	221	12	5,4%
Pharmacogenomics	231	231	100%

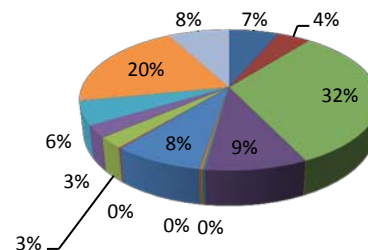
Disease risk results

Gen	Mutación	Enfermedad	Nº pacientes
APC	c.3920T>A, p.Ile1307Lys	175100. FAMILIAL ADENOMATOUS POLYPOSIS 1	2
ATP7B	c.1934T>G, p.Met645Arg	277900. WILSON DISEASE	10
BRCA2	c.658_659delGT p.Val220IlefsTer4	114480. BREAST CANCER	1
MUTYH	c.1187G>A p.Gly396Asp	608456. FAMILIAL ADENOMATOUS POLYPOSIS 2; FAP2	5
MUTYH	c.536A>G, p.Tyr179Cys	608456. FAMILIAL ADENOMATOUS POLYPOSIS 2; FAP2	2

PHARMGKB

Pharmacogenetic variants distribution

- CYP2C19 Clopidogrel, Citalopram/Escitalopram; Amitriptyline; Voriconazol
- CYP2C9 Fenitoína
- CYP2D6 Diversos fármacos: amitriptyline/ antidepressants/ clomipramine/ desipramine...
- CYP3A5 Tacrolimus
- DPYD Capecitabina, 5-FU, Tegafur
- G6PD Rasburicasa, Cloroquina, Dapsona, Clorpropamida.
- IFNL3 Peg-interferón y ribavirina
- NUDT15 Tiopurinas





Conclusions

- ▶ NAGEN 1000 as a model for Personalised genomic medicine implementation in the Public Health Service:
 - ▶ Multidisciplinary participation of professionals and patients.
 - ▶ Optimizes the use of resources and the development of new ones.
 - ▶ Provides training, specialization.
 - ▶ Produces scientific findings that increase knowledge.
 - ▶ Opens the integration and analysis of big data health and promotes innovative ICT solutions.
 - ▶ Encourages the adaptation of ethical legal principles to new technologies.
 - ▶ Facilitates cooperation between sectors.
 - ▶ Evidences the need for national and international strategies.
 - ▶ Enables the application of Genomic Medicine and prepares de basis of Personalised Medicine.

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PROYECTO 1000
GENOMA NAVARRA



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