

A decade of seascape genetics: contributions to basic and applied marine connectivity

Kimberly A. Selkoe*, Cassidy C. D'Aloia, Eric D. Crandall, Matthew Iacchei, Libby Liggins, Jonathan B. Puritz, Sophie von der Heyden, Robert J. Toonen

*Corresponding author: selkoe@nceas.ucsb.edu

Marine Ecology Progress Series 554: 1–19 (2016)

Supplementary Methods

Building a database of seascape genetic papers

To generate a database of seascape genetic papers, we conducted Web of Science searches in January 2016 using the search terms (seascape genetic*), (seascape genom*), (marine landscape population genetic*), and (SNP marine selection OR SNP marine adaptation) with the dates restricted from 2006 to 2015. We excluded papers that were: (1) clearly irrelevant to the topic; (2) review papers; (3) purely descriptive and lacked statistical analyses; and/or (4) did not use empirical genetic data. Next, we carefully read the papers and only retained those that directly tested for the influence of seascape features on gene flow, genetic structure, genetic diversity and/or connectivity. Finally, we supplemented the database with additional seascape genetic papers that were not indexed by Web of Science using our search terms, but were gathered by broad reading of the literature by the co-authors of this review. These papers were subject to the same filtering criteria listed above.

We acknowledge that the search terms used may have missed many studies of diadromous species. We retained all diadromous studies found, as we believe that seascape genetics can learn from these studies, which bridge the landscape-seascape divide. Overall, this process led to the inclusion of 100 seascape genetic papers in the database. We used all 100 papers to assess trends in marker usage (Fig. 3—main text) as well as the taxonomic and geographic breadth of studies (Fig. 4—main text).

Exploring seascape predictors

To explore common drivers of genetic structuring, we investigated a subset of our 100 papers that statistically analyzed the relationship between multiple seascape predictors and a genetic response metric. Here, we excluded papers that (1) only considered one predictor variable, or (2) only used simple Mantel tests, correlations, or simple linear regressions. This led to 53 studies whose trends are discussed in detail in the main text. These studies are identified in the reference list below with an asterisk.

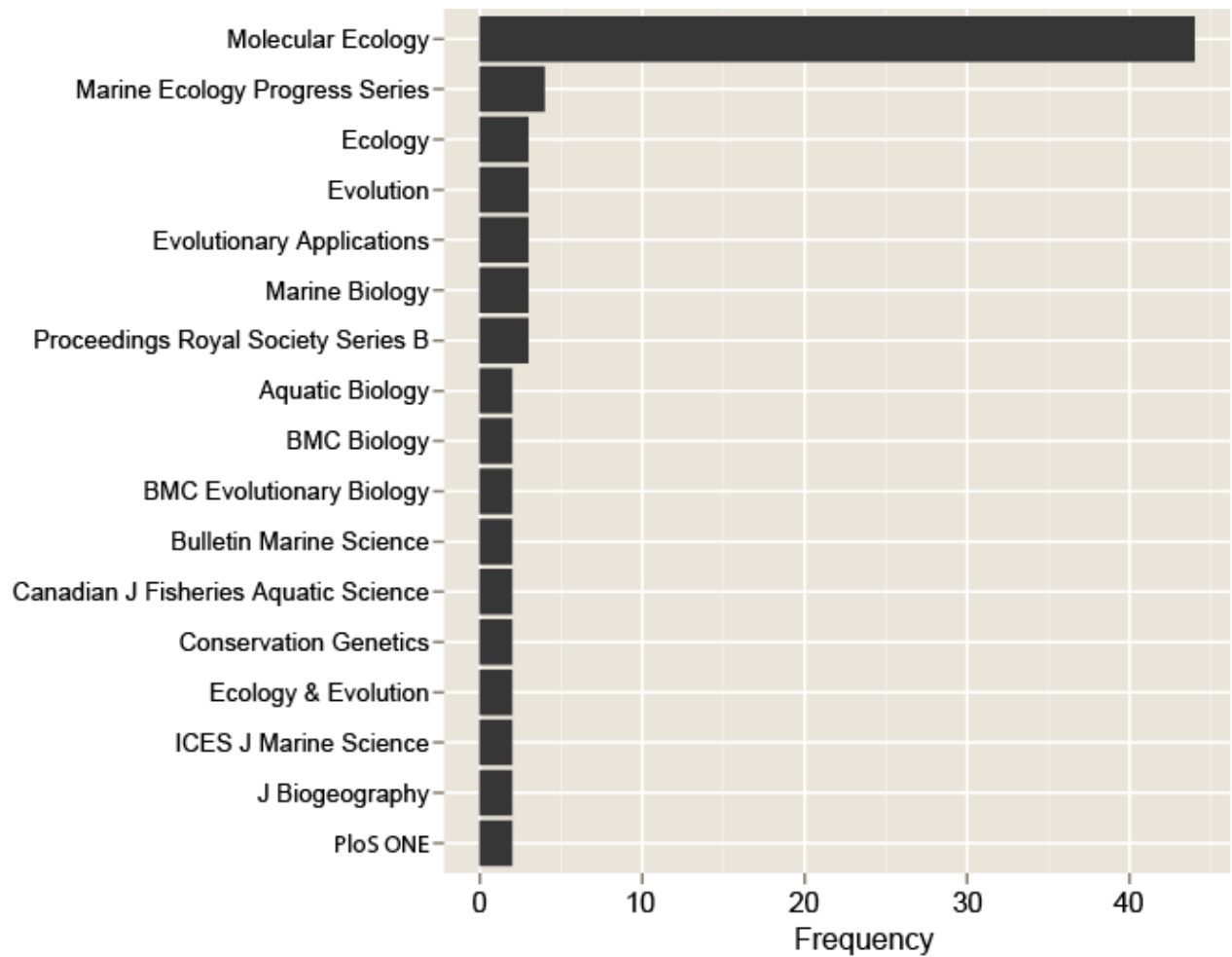


Fig S1. Seventeen journals that have published at least two seascape genetic papers during 2006-2015. Considering all journals that published at least one study, papers have appeared in 34 unique journals.

Table S1. Seascape genetic studies (n = 100) coded by: the journal they appear in, the focal taxonomic group, the marker used, ecosystem type, global position, and ocean/major sea. Data are also available as an Excel spreadsheet. Ang. is Angiospermae; Diadr. is diadromous; Estuar. is estuarine; GOM is Gulf of Mexico. Note that ‘Microsatellite’ refers to putatively neutral microsatellites; gene-associated microsatellites were classified as ‘Other’.

Year	First Author	Journal	Taxon	Marker	Ecosystem	Global Position	Oceans
2015	Berg	Genome Biol Evol	Chordata	SNPs	Subtidal	Temperate	North Sea/Baltic Sea
2015	Bradbury	Mol Ecol	Chordata	Multiple	Diadr. or Estuar.	Temperate	N. Atlantic Estuaries
2015	Chaves-Fonnegra	Mol Ecol	Porifera	Microsatellites	Subtidal	Tropical	Caribbean
2015	Cossu	Hydrobiologia	Mollusca	Multiple	Intertidal + subtidal	Temperate	Mediterranean
2015	Daloia	PNAS	Chordata	Microsatellites	Subtidal	Tropical	Caribbean
2015	Davies	Mol Ecol	Cnidaria	Microsatellites	Subtidal	Tropical	North Pacific
2015	Deli	Acta Zool Bulgar	Arthropoda	Other	Intertidal	Temperate	Mediterranean
2015	Giles	Ecol Evol	Porifera	Microsatellites	Subtidal	Tropical	Red Sea
2015	Gonçalves da Silva	Mol Ecol	Chordata	SNPs	Subtidal	Temperate	South Pacific
2015	Gorospe	PloS ONE	Cnidaria	Microsatellites	Subtidal	Tropical	North Pacific
2015	Guo	BMC Biology	Chordata	SNPs	Diadr. or Estuar.	Temperate	North Sea/Baltic Sea
2015	Hecht	Mol Ecol	Chordata	SNPs	Diadr. or Estuar.	Temperate	N. Pacific Estuaries
2015	Jahnke	Mol Ecol	Angiospermae	Microsatellites	Subtidal	Temperate	Mediterranean
2015	Johansson	Mol Ecol	Heterokontophyta	Microsatellites	Subtidal	Temperate	North Pacific
2015	Jorde	Mol Ecol	Arthropoda	Microsatellites	Subtidal	Polar/Subpolar	North Atlantic
2015	Liggins (online first)	J Biogeogr	Chordata	mtDNA	Subtidal	Tropical	Indo-Pacific
2015	Mege	Mar Ecol	Cnidaria	Microsatellites	Subtidal	Tropical	Caribbean
2015	Portnoy	Mol Ecol	Chordata	Multiple	Subtidal	Temperate	N. Atlantic + GOM
2015	Quemere	J Biogeogr	Chordata	Microsatellites	Diadr. or Estuar.	Temperate	N. Atlantic Estuaries
2015	Saenz-Agudelo	Mol Ecol	Chordata	SNPs	Subtidal	Tropical	Red Sea
2015	Saha	ICES J Mar Sci	Chordata	SNPs	Subtidal	Temperate	North Atlantic
2015	Sandoval-Castillo	MEPS	Chordata	Multiple	Subtidal	Temperate	North Pacific
2015	Sjoqvist	Mol Ecol	Heterokontophyta	Microsatellites	Pelagic	Temperate	North Sea/Baltic Sea
2015	Teske	MEPS	Mollusca	Microsatellites	Intertidal	Temperate	Indian + S. Pacific
2015	Thomas	Proc Roy Soc B	Cnidaria	Microsatellites	Subtidal	Tropical	Indian
2015	Young	Evol Appl	Chordata	Microsatellites	Subtidal	Polar/Subpolar	Southern
2014	Crandall	Bull Mar Sci	Echinodermata	mtDNA	Subtidal	Tropical	Indo-Pacific
2014	Daloia	Mol Ecol	Chordata	Multiple	Subtidal	Tropical	Caribbean
2014	Kesaniemi	Mar Biol	Annelida	Microsatellites	Diadr. or Estuar.	Temperate	North Sea/Baltic Sea
2014	Liggins	Bull Mar Sci	Echinodermata	mtDNA	Subtidal	Tropical	Indo-Pacific
2014	Milano	Mol Ecol	Chordata	SNPs	Subtidal	Temperate	N. Atlantic + Mediterr.

2014	Munguia-Vega	PeerJ	Chordata	mtDNA	Subtidal	Temperate	Gulf of California
2014	Nakajima	Mol Ecol	Angiospermae	Microsatellites	Intertidal + subtidal	Tropical	North Pacific
2014	Nanninga	Mol Ecol	Chordata	Microsatellites	Subtidal	Tropical	Red Sea
2014	Oliva	Mol Ecol	Angiospermae	Microsatellites	Subtidal	Temperate	Mediterranean
2014	Pujolar	Mol Ecol	Chordata	SNPs	Diadr. or Estuar.	Temperate	N. Atlantic Estuaries
2014	Robuchon	Mol Ecol	Heterokontophyta	Microsatellites	Intertidal + subtidal	Temperate	North Atlantic
2014	Rouger	Mol Ecol	Angiospermae	Microsatellites	Intertidal	Temperate	North Atlantic
2014	Schiavina	Mol Ecol	Arthropoda	Microsatellites	Intertidal	Temperate	Mediterranean
2014	Serrano	Mol Ecol	Cnidaria	Microsatellites	Subtidal	Tropical	Caribbean
2014	Sunday	Mol Ecol	Echinodermata	Microsatellites	Subtidal	Temperate	North Pacific
2014	Vandamme	Mol Ecol	Chordata	Multiple	Subtidal	Temperate	North Atlantic
2014	Viricel	Mol Ecol	Chordata	Multiple	Pelagic	Tropical	North Atlantic
2013	Andras	Mol Ecol	Cnidaria	Microsatellites	Subtidal	Tropical	Caribbean
2013	Bradbury	CJFAS	Chordata	Microsatellites	Diadr. or Estuar.	Temperate	N. Atlantic Estuaries
2013	Chust	Estuar Coast Shelf	Ang. + Mollusca	Microsatellites	Diadr. or Estuar.	Temperate	North Atlantic
2013	Coscia	Conserv Genet	Mollusca	Microsatellites	Intertidal	Temperate	North Atlantic
2013	DeFaveri	Evolution	Chordata	Multiple	Diadr. or Estuar.	Temperate	North Sea/Baltic Sea
2013	Gkafas	Aquat Biol	Chordata	Microsatellites	Subtidal	Temperate	Mediterranean
2013	Godhe	Proc Roy Soc B	Heterokontophyta	Microsatellites	Subtidal	Temperate	North Sea/Baltic Sea
2013	Hasselmann	Mol Ecol	Chordata	Microsatellites	Diadr. or Estuar.	Temperate	N. Atlantic Estuaries
2013	Hess	Mol Ecol	Chordata	SNPs	Diadr. or Estuar.	Temperate	N. Pacific Estuaries
2013	Iacchei	Mol Ecol	Arthropoda	Multiple	Subtidal	Temperate	North Pacific
2013	Krueger-Hadfield	Mol Ecol	Rhodophyta	Microsatellites	Intertidal	Temperate	North Atlantic
2013	Lee	J R Soc Interface	Cnidaria	Multiple	Pelagic	Temperate	North Atlantic
2013	Pespeni	Mol Ecol	Echinodermata	Other	Intertidal + subtidal	Temperate	North Pacific
2013	Saavedra-Sotelo	Ecol Evol	Cnidaria	Other	Subtidal	Tropical	North Pacific
2013	Teacher	Evol Appl	Chordata	Other	Pelagic	Temperate	North Sea/Baltic Sea
2013	Therkildsen	Evol Appl	Chordata	SNPs	Subtidal	Polar/Subpolar	North Atlantic
2013	Vincent	Evolution	Chordata	SNPs	Diadr. or Estuar.	Temperate	N. Atlantic Estuaries
2013	Wei	MEPS	Mollusca	Microsatellites	Intertidal + subtidal	Temperate	South Pacific
2013	Wei	Mar Biol	Mollusca	Microsatellites	Intertidal + subtidal	Temperate	South Pacific
2012	Amaral	PloS ONE	Chordata	Microsatellites	Pelagic	Temperate	Multiple Oceans
2012	Berry	Mol Ecol	Chordata	Microsatellites	Subtidal	Tropical	Indian + S. Pacific
2012	Berry	Fish Oceanogr	Chordata	Microsatellites	Subtidal	Tropical	Indian
2012	Crandall	Mol Ecol	Mollusca	mtDNA	Intertidal + Diadr.	Tropical	South Pacific

2012	Foster	Mol Ecol	Cnidaria	Microsatellites	Subtidal	Tropical	Caribbean
2012	Knutsen	MEPS	Chordata	Microsatellites	Subtidal	Temperate	North Atlantic
2012	Limborg	Mol Ecol	Chordata	SNPs	Subtidal	Temperate	North Atlantic
2012	Paz-Garcia	Pac Sci	Cnidaria	Other	Subtidal	Tropical	North Pacific
2012	Roy	CJFAS	Chordata	Microsatellites	Subtidal	Temperate	North Atlantic
2011	Alberto	Mol Ecol	Heterokontophyta	Microsatellites	Subtidal	Temperate	North Pacific
2011	Coleman	J Ecol	Heterokontophyta	Microsatellites	Subtidal	Temperate	South Pacific + Indian
2011	Matala	T Am Fish Soc	Chordata	Multiple	Diadr. or Estuar.	Temperate	N. Pacific Estuaries
2011	Mendez	Heredity	Chordata	mtDNA	Subtidal	Tropical	Indian
2011	Olsen	Conserv Genet	Chordata	Microsatellites	Diadr. or Estuar.	Polar/Subpolar	N. Pacific Estuaries
2011	Puritz	Nature Commun	Echinodermata	Multiple	Intertidal + subtidal	Temperate	North Pacific
2011	Schunter	Mol Ecol	Chordata	Microsatellites	Subtidal	Temperate	Mediterranean
2010	Alberto	Ecology	Heterokontophyta	Microsatellites	Subtidal	Temperate	North Pacific
2010	Banks	Mol Ecol	Echinodermata	Microsatellites	Subtidal	Temperate	South Atlantic
2010	Fraser	BMC Evol Biol	Heterokontophyta	Multiple	Subtidal	Temperate	South Pacific
2010	Galindo	Mol Ecol	Arthropoda	Multiple	Intertidal	Temperate	North Pacific
2010	Gonzalez-Wanguemert	J Exp Mar Biol Ecol	Chordata	Multiple	Subtidal	Temperate	N. Atlantic + Mediterr.
2010	Mendez	Mol Ecol	Chordata	Multiple	Subtidal	Temperate	South Atlantic
2010	Selkoe	Mol Ecol	Multiple	Microsatellites	Subtidal	Temperate	North Pacific
2010	White	Proc Roy Soc B	Mollusca	Microsatellites	Subtidal	Temperate	North Pacific
2010	White	Mol Ecol	Chordata	Microsatellites	Subtidal	Temperate	North Atlantic
2009	Gaggiotti	Evolution	Chordata	Microsatellites	Pelagic	Temperate	North Sea/Baltic Sea
2009	Gonzalez-Wanguemert	Biochem Syst Ecol	Mollusca	Other	Subtidal	Temperate	Mediterranean
2009	Knutsen	Mol Ecol	Chordata	Microsatellites	Subtidal	Polar/Subpolar	North Atlantic
2009	McInerney	Aquat Biol	Mollusca	Microsatellites	Intertidal	Temperate	North Atlantic
2009	Nielsen	BMC Evol Biol	Chordata	SNPs	Subtidal	Temperate	North Atlantic
2009	Pampoulie	ICES J Mar Sci	Chordata	Microsatellites	Subtidal	Polar/Subpolar	North Atlantic
2009	Sala-Bozano	Mol Ecol	Chordata	Multiple	Subtidal	Temperate	N. Atlantic + Mediterr.
2008	Schultz	Mol Ecol	Chordata	Multiple	Subtidal	Tropical	Multiple Oceans
2007	Banks	Ecology	Echinodermata	Microsatellites	Subtidal	Temperate	South Pacific
2007	Fontaine	BMC Biology	Chordata	Microsatellites	Subtidal	Temperate	North Atlantic
2006	Barreiro	Mar Biol	Mollusca	Other	Subtidal	Temperate	North Atlantic
2006	Galindo	Curr Biol	Cnidaria	Multiple	Subtidal	Tropical	Caribbean
2006	Selkoe	Ecology	Chordata	Microsatellites	Subtidal	Temperate	North Pacific

Reference list for seascape genetic papers from 2006-2015

- *Papers included in the analyses of multiple seascape predictors are denoted by an asterisk.
- *Alberto F, Raimondi PT, Reed DC, Coelho NC, Leblois R, Whitmer A, Serrão EA (2010) Habitat continuity and geographic distance predict population genetic differentiation in giant kelp. *Ecology* 91:49–56
- Alberto F, Raimondi PT, Reed DC, Watson JR, Siegel DA, Mitarai S, Coelho N, Serrão EA (2011) Isolation by oceanographic distance explains genetic structure for *Macrocystis pyrifera* in the Santa Barbara Channel. *Mol Ecol* 20:2543–2554
- *Amaral AR, Beheregaray LB, Bilgmann K, Boutov D, Freitas L, Robertson KM, Sequeira M, Stockin KA, Coelho MM, Möller LM (2012) Seascape genetics of a globally distributed, highly mobile marine mammal: the short-beaked common dolphin (genus *Delphinus*). *PLoS One* 7:e31482
- Andras JP, Rypien KL, Harvell CD (2013) Range-wide population genetic structure of the Caribbean sea fan coral, *Gorgonia ventalina*. *Mol Ecol* 22:56–73
- Banks SC, Piggott MP, Williamson JE, Bové U, Holbrook NJ, Beheregaray LB (2007) Oceanic variability and coastal topography shape genetic structure in a long-dispersing sea urchin. *Ecology* 88:3055–3064
- *Banks SC, Ling SD, Johnson CR, Piggott MP, Williamson JE, Beheregaray LB (2010) Genetic structure of a recent climate change-driven range extension. *Mol Ecol* 19:2011–2024
- Barreiro R, Couceiro L, Quintela M, Ruiz JM (2006) Population genetic structure of the prosobranch *Nassarius reticulatus* (L.) in a ria seascape (NW Iberian Peninsula) as revealed by RAPD analysis. *Mar Biol* 148:1051–1060
- *Berg PR, Jentoft S, Star B, Ring KH, Knutsen H, Lien S, Jakobsen KS, André C (2015) Adaptation to low salinity promotes genomic divergence in Atlantic cod (*Gadus morhua* L.). *Genome Biol Evol* 7:1644–1663
- Berry O, England P, Fairclough D, Jackson G, Greenwood JIM (2012a) Microsatellite DNA analysis and hydrodynamic modelling reveal the extent of larval transport and gene flow between management zones in an exploited marine fish (*Glaucosoma hebraicum*). *Fish Oceanogr* 21:243–254
- Berry O, England P, Marriott RJ, Burrridge CP, Newman SJ (2012b) Understanding age-specific dispersal in fishes through hydrodynamic modelling, genetic simulations and microsatellite DNA analysis. *Mol Ecol* 21:2145–2159
- *Bradbury IR, Hamilton LC, Robertson MJ, Bourgeois CE, Mansour A, Dempson JB, Marshall CT (2013) Landscape structure and climatic variation determine Atlantic salmon genetic connectivity in the Northwest Atlantic. *Can J Fish Aquat Sci* 71:246–258
- Bradbury IR, Hamilton LC, Dempson B, Robertson MJ, Bourret V, Bernatchez L, Verspoor E (2015) Transatlantic secondary contact in Atlantic Salmon, comparing microsatellites, a single nucleotide polymorphism array and restriction-site associated DNA sequencing for the resolution of complex spatial structure. *Mol Ecol* 24:5130–5144
- Chaves-Fonnegra A, Feldheim KA, Secord J, Lopez JV (2015) Population structure and dispersal of the coral-excavating sponge *Cliona delitrix*. *Mol Ecol* 24:1447–1466
- Chust G, Albaina A, Aranburu A, Borja Á, Diekmann OE, Estonba A, Franco J, Garmendia JM, Iriondo M, Muxika I (2013) Connectivity, neutral theories and the assessment of species vulnerability to global change in temperate estuaries. *Estuar Coast Shelf S* 131:52–63
- Coleman MA, Roughan M, Macdonald HS, Connell SD, Gillanders BM, Kelaher BP, Steinberg PD (2011) Variation in the strength of continental boundary currents determines continent-wide connectivity in kelp. *J Ecol* 99:1026–1032

- Coscia I, Robins PE, Porter JS, Malham SK, Ironside JE (2013) Modelled larval dispersal and measured gene flow: seascape genetics of the common cockle *Cerastoderma edule* in the southern Irish Sea. *Conserv Genet* 14:451–466
- *Cossu P, Dedola GL, Scarpa F, Sanna D, Lai T, Maltagliati F, Curini-Galletti M, Casu M (2015) Patterns of spatial genetic variation in *Patella ulyssiponensis*: insights from the western Mediterranean marine ecoregion. *Hydrobiologia* 755:39–55
- *Crandall ED, Trembl EA, Barber PH (2012) Coalescent and biophysical models of stepping-stone gene flow in neritid snails. *Mol Ecol* 21:5579–5598
- *Crandall ED, Trembl EA, Liggins L, Gleeson L, Yasuda N, Barber PH, Wörheide G, Riginos C (2014) Return of the ghosts of dispersal past: historical spread and contemporary gene flow in the blue sea star *Linckia laevigata*. *B Mar Sci* 90:399–425
- *D'Aloia CC, Bogdanowicz SM, Harrison RG, Buston PM (2014) Seascape continuity plays an important role in determining patterns of spatial genetic structure in a coral reef fish. *Mol Ecol* 23:2902–2913
- D'Aloia CC, Bogdanowicz SM, Francis RK, Majoris JE, Harrison RG, Buston PM (2015) Patterns, causes, and consequences of marine larval dispersal. *P Natl Acad Sci USA* 112:13940–13945
- Davies SW, Trembl EA, Kenkel CD, Matz MV (2015) Exploring the role of Micronesian islands in the maintenance of coral genetic diversity in the Pacific Ocean. *Mol Ecol* 24:70–82
- *DeFaveri J, Jonsson PR, Merilä J (2013) Heterogeneous genomic differentiation in marine threespine sticklebacks: adaptation along an environmental gradient. *Evolution* 67:2530–2546
- Deli T, Said K, Chatti N (2015) Genetic Differentiation among Populations of the Green Crab *Carcinus aestuarii* (Nardo, 1847) (Brachyura, Carcinidae) from the Eastern and Western Mediterranean Coast of Tunisia. *Acta Zool Bulgar* 67:327–335
- Fontaine MC, Baird SJ, Piry S, Ray N, Tolley KA, Duke S, Birkun A, Ferreira M, Jauniaux T, Llavona Á (2007) Rise of oceanographic barriers in continuous populations of a cetacean: the genetic structure of harbour porpoises in Old World waters. *BMC Biol* 5:1
- Foster NL, Paris CB, Kool JT, Baums IB, Stevens JR, Sanchez JA, Bastidas C, Agudelo C, Bush P, Day O (2012) Connectivity of Caribbean coral populations: complementary insights from empirical and modelled gene flow. *Mol Ecol* 21:1143–1157
- *Fraser CI, Thiel M, Spencer HG, Waters JM (2010) Contemporary habitat discontinuity and historic glacial ice drive genetic divergence in Chilean kelp. *BMC Evol Biol* 10:203
- Gaggiotti OE, Bekkevold D, Jørgensen HB, Foll M, Carvalho GR, Andre C, Ruzzante DE (2009) Disentangling the effects of evolutionary, demographic, and environmental factors influencing genetic structure of natural populations: Atlantic herring as a case study. *Evolution* 63:2939–2951
- *Galindo HM, Olson DB, Palumbi SR (2006) Seascape genetics: a coupled oceanographic-genetic model predicts population structure of Caribbean corals. *Curr Biol* 16:1622–1626
- Galindo HM, Pfeiffer-Herbert AS, McManus MA, Chao YI, Chai FEI, Palumbi SR (2010) Seascape genetics along a steep cline: using genetic patterns to test predictions of marine larval dispersal. *Mol Ecol* 19:3692–3707
- *Giles EC, Saenz-Agudelo P, Hussey NE, Ravasi T, Berumen ML (2015) Exploring seascape genetics and kinship in the reef sponge *Stylissa carteri* in the Red Sea. *Ecol Evol* 5:2487–2502
- Gkafas GA, Tsigenopoulos C, Magoulas A, Panagiotaki P, Vafidis D, Mamuris Z, Exadactylos A (2013) Population subdivision of saddled seabream *Oblada melanura* in the Aegean Sea revealed by genetic and morphometric analyses. *Aquat Biol* 18:69–80

- Godhe A, Egardt J, Kleinhans D, Sundqvist L, Hordoir R, Jonsson PR (2013) Seascape analysis reveals regional gene flow patterns among populations of a marine planktonic diatom. *P Roy Soc B- Biol Sci* 280:20131599
- Gonçalves da Silva A, Appleyard SA, Upston J (2015) Establishing the evolutionary compatibility of potential sources of colonizers for overfished stocks: a population genomics approach. *Mol Ecol* 24:564–579
- *González-Wangüemert M, Cánovas F, Marcos C, Pérez-Ruzafa Á (2009) Phosphoglucose isomerase variability of *Cerastoderma glaucum* as a model for testing the influence of environmental conditions and dispersal patterns through quantitative ecology approaches. *Biochem Syst Ecol* 37:325–333
- González-Wangüemert M, Cánovas F, Pérez-Ruzafa A, Marcos C, Alexandrino P (2010) Connectivity patterns inferred from the genetic structure of white seabream (*Diplodus sargus* L.). *J Exp Mar Biol Ecol* 383:23–31
- *Goroppe KD, Karl SA (2015) Depth as an organizing force in *Pocillopora damicornis*: intra-reef genetic architecture. *PloS one* 10:e0122127
- *Guo B, DeFaveri J, Sotelo G, Nair A, Merilä J (2015) Population genomic evidence for adaptive differentiation in Baltic Sea three-spined sticklebacks. *BMC Biol* 13:1
- *Hasselman DJ, Ricard D, Bentzen P (2013) Genetic diversity and differentiation in a wide ranging anadromous fish, American shad (*Alosa sapidissima*), is correlated with latitude. *Mol Ecol* 22:1558–1573
- *Hecht BC, Matala AP, Hess JE, Narum SR (2015) Environmental adaptation in Chinook salmon (*Oncorhynchus tshawytscha*) throughout their North American range. *Mol Ecol* 24:5573–5595
- Hess JE, Campbell NR, Close DA, Docker MF, Narum SR (2013) Population genomics of Pacific lamprey: adaptive variation in a highly dispersive species. *Mol Ecol* 22:2898–2916
- *Iacchei M, Ben-Horin T, Selkoe KA, Bird CE, García-Rodríguez FJ, Toonen RJ (2013) Combined analyses of kinship and FST suggest potential drivers of chaotic genetic patchiness in high gene-flow populations. *Mol Ecol* 22:3476–3494
- *Jahnke M, Olsen JL, Procaccini G (2015) A meta-analysis reveals a positive correlation between genetic diversity metrics and environmental status in the long-lived seagrass *Posidonia oceanica*. *Mol Ecol* 24:2336–2348
- *Johansson ML, Alberto F, Reed DC, Raimondi PT, Coelho NC, Young MA, Drake PT, Edwards CA, Cavanaugh K, Assis J (2015) Seascape drivers of *Macrocystis pyrifera* population genetic structure in the northeast Pacific. *Mol Ecol* 24:4866–4885
- *Jorde PE, Søvik G, Westgaard J-I, Albretsen J, André C, Hvingel C, Johansen T, Sandvik AD, Kingsley M, Jørstad KE (2015) Genetically distinct populations of northern shrimp, *Pandalus borealis*, in the North Atlantic: adaptation to different temperatures as an isolation factor. *Mol Ecol* 24:1742–1757
- *Kesäniemi JE, Hansen BW, Banta GT, Knott KE (2014) Chaotic genetic patchiness and high relatedness of a poecilogonous polychaete in a heterogeneous estuarine landscape. *Mar Biol* 161:2631–2644
- *Knutsen H, Jorde PE, Sannaes H, Rus Hoelzel A, Bergstad OA, Stefanni S, Johansen T, Stenseth NC (2009) Bathymetric barriers promoting genetic structure in the deepwater demersal fish tusk (*Brosme brosme*). *Mol Ecol* 18:3151–3162
- Knutsen H, Jorde PE, Bergstad OA, Skogen M (2012) Population genetic structure in a deepwater fish *Coryphaenoides rupestris*: patterns and processes. *Mar Ecol Prog Ser* 460:233–246

- *Krueger-Hadfield SA, Roze D, Mauger S, Valero M (2013) Intergametophytic selfing and microgeographic genetic structure shape populations of the intertidal red seaweed *Chondrus crispus*. *Mol Ecol* 22:3242–3260
- *Lee PL, Dawson MN, Neill SP, Robins PE, Houghton JD, Doyle TK, Hays GC (2013) Identification of genetically and oceanographically distinct blooms of jellyfish. *J R Soc Interface* 10:20120920
- Liggins L, Gleeson L, Riginos C (2014) Evaluating edge-of-range genetic patterns for tropical echinoderms, *Acanthaster planci* and *Tripneustes gratilla*, of the Kermadec Islands, southwest Pacific. *B Mari Sci* 90:379–397
- *Liggins L, Treml EA, Possingham HP, Riginos C (2016)[†] Seascape features, rather than dispersal traits, predict spatial genetic patterns in co-distributed reef fishes. *J Biogeogr* 43:256–267
- [†]Online first edition from 2015 counted in database
- *Limborg MT, Helyar SJ, Bruyn M de, Taylor MI, Nielsen EE, Ogden ROB, Carvalho GR, Bekkevold D (2012) Environmental selection on transcriptome-derived SNPs in a high gene flow marine fish, the Atlantic herring (*Clupea harengus*). *Mol Ecol* 21:3686–3703
- *Matala AP, Hess JE, Narum SR (2011) Resolving adaptive and demographic divergence among Chinook salmon populations in the Columbia River basin. *T Am Fish Soc* 140:783–807
- McInerney CE, Allcock AL, Johnson MP, Prodöhl PA (2009) Understanding marine reserve function in a seascape genetics context: *Nucella lapillus* in Strangford Lough (Northern Ireland) as an example. *Aquat Biol* 7:45–58
- Mège P, Schizas NV, Garcia Reyes J, Hrbek T (2015) Genetic seascape of the threatened Caribbean elkhorn coral, *Acropora palmata*, on the Puerto Rico Shelf. *Mar Ecol* 36:195–209
- *Mendez M, Rosenbaum HC, Subramaniam A, Yackulic C, Bordino P (2010) Isolation by environmental distance in mobile marine species: molecular ecology of franciscana dolphins at their southern range. *Mol Ecol* 19:2212–2228
- *Mendez M, Subramaniam A, Collins T, Minton G, Baldwin R, Berggren P, Särnblad A, Amir OA, Peddemors VM, Karczmarski L (2011) Molecular ecology meets remote sensing: environmental drivers to population structure of humpback dolphins in the Western Indian Ocean. *Heredity* 107:349–361
- *Milano I, Babbucci M, Cariani A, Atanassova M, Bekkevold D, Carvalho GR, Espiñeira M, Fiorentino F, Garofalo G, Geffen AJ (2014) Outlier SNP markers reveal fine-scale genetic structuring across European hake populations (*Merluccius merluccius*). *Mol Ecol* 23:118–135
- Munguia-Vega A, Jackson A, Marinone SG, Erisman B, Moreno-Baez M, Girón-Nava A, Pfister T, Aburto-Oropeza O, Torre J (2014) Asymmetric connectivity of spawning aggregations of a commercially important marine fish using a multidisciplinary approach. *PeerJ* 2:e511
- Nakajima Y, Matsuki Y, Lian C, Fortes MD, Uy WH, Campos WL, Nakaoka M, Nadaoka K (2014) The Kuroshio Current influences genetic diversity and population genetic structure of a tropical seagrass, *Enhalus acoroides*. *Mol Ecol* 23:6029–6044
- *Nanninga GB, Saenz-Agudelo P, Manica A, Berumen ML (2014) Environmental gradients predict the genetic population structure of a coral reef fish in the Red Sea. *Mol Ecol* 23:591–602
- *Nielsen EE, Hemmer-Hansen J, Poulsen NA, Loeschcke V, Moen T, Johansen T, Mittelholzer C, Taranger G-L, Ogden R, Carvalho GR (2009) Genomic signatures of local directional selection in a high gene flow marine organism; the Atlantic cod (*Gadus morhua*). *BMC Evol Biol* 9:276
- *Oliva S, Romero J, Pérez M, Manent P, Mascaró O, Serrão EA, Coelho N, Alberto F (2014) Reproductive strategies and isolation-by-demography in a marine clonal plant along an eutrophication gradient. *Mol Ecol* 23:5698–5711

- *Olsen JB, Crane PA, Flannery BG, Dunmall K, Templin WD, Wenburg JK (2011) Comparative landscape genetic analysis of three Pacific salmon species from subarctic North America. *Conserv Genet* 12:223–241
- Pampoulie C, Gíslason D, Daniélsdóttir AK (2009) A “seascape genetic” snapshot of *Sebastes marinus* calls for further investigation across the North Atlantic. *ICES J Mar Sci* 66:2219–2222
- Paz-García DA, Chávez-Romo HE, Correa-Sandoval F, Reyes-Bonilla H, López-Pérez A, Medina-Rosas P, Hernández-Cortés MP (2012) Genetic connectivity patterns of corals *Pocillopora damicornis* and *Porites panamensis* (Anthozoa: Scleractinia) along the west coast of Mexico. *Pac Sci* 66:43–61
- *Pespeni MH, Palumbi SR (2013) Signals of selection in outlier loci in a widely dispersing species across an environmental mosaic. *Mol Ecol* 22:3580–3597
- *Portnoy DS, Puritz JB, Hollenbeck CM, Gelsleichter J, Chapman D, Gold JR (2015) Selection and sex-biased dispersal in a coastal shark: the influence of philopatry on adaptive variation. *Mol Ecol* 24:5877–5885
- *Pujolar JM, Jacobsen MW, Als TD, Frydenberg J, Munch K, Jónsson B, Jian JB, Cheng L, Maes GE, Bernatchez L (2014) Genome-wide single-generation signatures of local selection in the panmictic European eel. *Mol Ecol* 23:2514–2528
- *Puritz JB, Toonen RJ (2011) Coastal pollution limits pelagic larval dispersal. *Nat Commun* 2:226
- *Quéméré E, Baglinière J-L, Roussel J-M, Evanno G, McGinnity P, Launey S (2015) Seascape and its effect on migratory life-history strategy influences gene flow among coastal brown trout (*Salmo trutta*) populations in the English Channel. *J Biogeography*
- Robuchon M, Le Gall L, Mauger S, Valero M (2014) Contrasting genetic diversity patterns in two sister kelp species co-distributed along the coast of Brittany, France. *Mol Ecol* 23:2669–2685
- *Rouger R, Jump AS (2014) A seascape genetic analysis reveals strong biogeographical structuring driven by contrasting processes in the polyploid saltmarsh species *Puccinellia maritima* and *Triglochin maritima*. *Mol Ecol* 23:3158–3170
- *Roy D, Hurlbut TR, Ruzzante DE, Fraser DJ (2012) Biocomplexity in a demersal exploited fish, white hake (*Urophycis tenuis*): depth-related structure and inadequacy of current management approaches. *Can J Fish Aquat Sci* 69:415–429
- *Saavedra-Sotelo NC, Calderon-Aguilera LE, Reyes-Bonilla H, Paz-García DA, López-Pérez RA, Cupul-Magaña A, Cruz-Barraza JA, Rocha-Olivares A (2013) Testing the genetic predictions of a biogeographical model in a dominant endemic Eastern Pacific coral (*Porites panamensis*) using a genetic seascape approach. *Ecol Evol* 3:4070–4091
- *Saenz-Agudelo P, Dibattista JD, Piatek MJ, Gaither MR, Harrison HB, Nanninga GB, Berumen ML (2015) Seascape genetics along environmental gradients in the Arabian Peninsula: insights from ddRAD sequencing of anemonefishes. *Mol Ecol* 24:6241–6255
- Saha A, Hauser L, Kent M, Planque B, Neat F, Kirubakaran TG, Huse I, Homrum EÍ, Fevolden S-E, Lien S (2015) Seascape genetics of saithe (*Pollachius virens*) across the North Atlantic using single nucleotide polymorphisms. *ICES J Mar Sci* 72:2732–2741
- Sala-Bozano M, Ketmaier V, Mariani S (2009) Contrasting signals from multiple markers illuminate population connectivity in a marine fish. *Mol Ecol* 18:4811–4826
- *Sandoval-Castillo J, Beheregaray LB (2015) Metapopulation structure informs conservation management in a heavily exploited coastal shark (*Mustelus henlei*). *Mar Ecol Prog Ser* 533:191

- Schiavina M, Marino IAM, Zane L, Melià P (2014) Matching oceanography and genetics at the basin scale. Seascape connectivity of the Mediterranean shore crab in the Adriatic Sea. *Mol Ecol* 23:5496–5507
- *Schultz JK, Feldheim KA, Gruber SH, Ashley MV, McGovern TM, Bowen BW (2008) Global phylogeography and seascape genetics of the lemon sharks (genus *Negaprion*). *Mol Ecol* 17:5336–5348
- Schunter C, Carreras-Carbonell J, Macpherson E, Tintoré J, Vidal-Vijande E, Pascual A, Guidetti P, Pascual M (2011) Matching genetics with oceanography: directional gene flow in a Mediterranean fish species. *Mol Ecol* 20:5167–5181
- *Selkoe KA, Gaines SD, Caselle JE, Warner RR (2006) Current shifts and kin aggregation explain genetic patchiness in fish recruits. *Ecology* 87:3082–3094
- *Selkoe KA, Watson JR, White C, Horin TB, Iacchei M, Mitarai S, Siegel DA, Gaines SD, Toonen RJ (2010) Taking the chaos out of genetic patchiness: seascape genetics reveals ecological and oceanographic drivers of genetic patterns in three temperate reef species. *Mol Ecol* 19:3708–3726
- Serrano X, Baums IB, O'Reilly K, Smith TB, Jones RJ, Shearer TL, Nunes FLD, Baker AC (2014) Geographic differences in vertical connectivity in the Caribbean coral *Montastraea cavernosa* despite high levels of horizontal connectivity at shallow depths. *Mol Ecol* 23:4226–4240
- Sjöqvist C, Godhe A, Jonsson PR, Sundqvist L, Kremp A (2015) Local adaptation and oceanographic connectivity patterns explain genetic differentiation of a marine diatom across the North Sea–Baltic Sea salinity gradient. *Mol Ecol* 24:2871–2885
- Sunday JM, Popovic I, Palen WJ, Foreman MGG, Hart MW (2014) Ocean circulation model predicts high genetic structure observed in a long-lived pelagic developer. *Mol Ecol* 23:5036–5047
- *Teacher AG, André C, Jonsson PR, Merilä J (2013) Oceanographic connectivity and environmental correlates of genetic structuring in Atlantic herring in the Baltic Sea. *Evolutionary Applications* 6:549–567
- *Teske PR, Sandoval-Castillo J, Sebille E van, Waters J, Beheregaray LB (2015) On-shelf larval retention limits population connectivity in a coastal broadcast spawner. *Mar Ecol Prog Ser* 532:1
- Therkildsen NO, Hemmer-Hansen J, Hedeholm RB, Wisz MS, Pampoulie C, Meldrup D, Bonanomi S, Retzel A, Olsen SM, Nielsen EE (2013) Spatiotemporal SNP analysis reveals pronounced biocomplexity at the northern range margin of Atlantic cod *Gadus morhua*. *Evol Appl* 6:690–705
- Thomas L, Kennington WJ, Stat M, Wilkinson SP, Kool JT, Kendrick GA (2015) Isolation by resistance across a complex coral reef seascape. *P Roy Soc B- Biol Sci* 282:20151217
- *Vandamme SG, Maes GE, Raeymaekers JAM, Cottenie K, Imsland AK, Hellemans B, Lacroix G, Mac Aoidh E, Martinsohn JT, Martínez P (2014) Regional environmental pressure influences population differentiation in turbot (*Scophthalmus maximus*). *Mol Ecol* 23:618–636
- * Vincent B, Dionne M, Kent MP, Lien S, Bernatchez L (2013) Landscape genomics in Atlantic salmon (*Salmo salar*): searching for gene–environment interactions driving local adaptation. *Evolution* 67:3469–3487
- *Viricel A, Rosel PE (2014) Hierarchical population structure and habitat differences in a highly mobile marine species: the Atlantic spotted dolphin. *Mol Ecol* 23:5018–5035
- Wei K, Wood AR, Gardner JP (2013a) Population genetic variation in the New Zealand greenshell mussel: locus-dependent conflicting signals of weak structure and high gene flow balanced against pronounced structure and high self-recruitment. *Mar Biol* 160:931–949

- *Wei K, Wood AR, Gardner JP (2013b) Seascape genetics of the New Zealand greenshell mussel: sea surface temperature explains macrogeographic scale genetic variation. *Mar Ecol Prog Ser* 477:107–121
- *White C, Selkoe KA, Watson J, Siegel DA, Zacherl DC, Toonen RJ (2010) Ocean currents help explain population genetic structure. *P Roy Soc B- Biol Sci* 277:1685–1694.
- *White TA, Stamford J, Rus Hoelzel A (2010) Local selection and population structure in a deep-sea fish, the roundnose grenadier (*Coryphaenoides rupestris*). *Mol Ecol* 19:216–226
- Young EF, Belchier M, Hauser L, Horsburgh GJ, Meredith MP, Murphy EJ, Pascoal S, Rock J, Tysklind N, Carvalho GR (2015) Oceanography and life history predict contrasting genetic population structure in two Antarctic fish species. *Evol Appl* 8:486–509